

09/625,191 SEQUENCE LISTING

	2111	8 11				36	QUEIN	CE L	1311	NG							
<110	>	Comm	uni,	Did	ier												
<120	>	COMP	OSIT:	IONS	AND	METI	HODS	СОМ	PRIS	ING	G-PR	OTEI	V CO	UPLE	D REC	EPTORS	5
<130	>	9409	/213:	2													
<140 <141	>	us 10 2002	0/07! -02-	9,38 [,] 20	4												
<150: <151:		us 0: 2001			3												
<160	>	50															
<170	>	Pate	ntIn	ver	sion	3.1											
<210: <211: <212: <213:	> >	1 1356 DNA Homo	sap ⁻	iens													
<220: <221: <222: <223:	> >	CDS (1).	. (13!	56)													
<400: atg : Met :	gag	1 tcc Ser	tca Ser	ccc Pro 5	atc Ile	ccc Pro	cag Gln	tca Ser	tca Ser 10	ggg Gly	aac Asn	tct Ser	tcc ser	act Thr 15	ttg Leu		48
ggg a	agg Arg	gtc Val	cct Pro 20	caa Gln	acc Thr	cca Pro	ggt Gly	ccc Pro 25	tct Ser	act Thr	gcc Ala	agt Ser	ggg Gly 30	gtc Val	ccg Pro		96
gag (Glu	gtg Val	ggg Gly 35	cta Leu	cgg Arg	gat Asp	gtt Val	gct Ala 40	tcg Ser	gaa Glu	tct Ser	gtg Val	gcc Ala 45	ctc Leu	ttc Phe	ttc Phe	1	.44
atg (Met	ctc Leu 50	ctg Leu	ctg Leu	gac Asp	ttg Leu	act Thr 55	gct Ala	gtg Val	gct Ala	ggc Gly	aat Asn 60	gcc Ala	gct Ala	gtg Val	atg Met	1	.92
gcc (Ala v 65	gtg Val	atc Ile	gcc Ala	aag Lys	acg Thr 70	cct Pro	gcc Ala	ctc Leu	cga Arg	aaa Lys 75	ttt Phe	gtc Val	ttc Phe	gtc Val	ttc Phe 80	2	40
cac (His I	ctc Leu	tgc Cys	ctg Leu	gtg Val 85	gac Asp	ctg Leu	ctg Leu	gct Ala	gcc Ala 90	ctg Leu	acc Thr	ctc Leu	atg Met	ccc Pro 95	ctg Leu	2	88
gcc a Ala M	atg Met	ctc Leu	tcc Ser 100	agc Ser	tct Ser	gcc Ala	ctc Leu	ttt Phe 105	gac Asp	cac His	gcc Ala	ctc Leu	ttt Phe 110	ggg Gly	gag Glu	3	36
gtg (Val /	gcc Ala	tgc Cys 115	cgc Arg	ctc Leu	tac Tyr	ttg Leu	ttt Phe 120	ctg Leu	agc Ser	gtg Val	tgc Cys	ttt Phe 125	gtc Val	agc Ser	ctg Leu	3	84
gcc a Ala :	atc Ile	ctc Leu	tcg Ser	gtg Val	tca Ser	gcc Ala	atc Ile	aat Asn	val	gag Glu Page	Arg	tac Tyr	tat Tyr	tac Tyr	gta Val	4	32

gtc Val 145	cac His	ccc Pro	atg Met	cgc Arg	tac Tyr 150	gag Glu	gtg Val	cgc Arg	atg Met	acg Thr 155	ctg Leu	ggg Gly	ctg Leu	gtg val	gcc Ala 160	480
tct Ser	gtg val	ctg Leu	gtg val	ggt Gly 165	gtg Val	tgg Trp	gtg Val	aag Lys	gcc Ala 170	ttg Leu	gcc Ala	atg Met	gct Ala	tct Ser 175	gtg Val	528
cca Pro	gtg Val	ttg Leu	gga Gly 180	agg Arg	gtc Val	tcc Ser	tgg Trp	gag Glu 185	gaa Glu	gga Gly	gct Ala	ccc Pro	agt Ser 190	gtc Val	ccc Pro	576
cca Pro	ggc Gly	tgt Cys 195	tca Ser	ctc Leu	cag Gln	tgg Trp	agc Ser 200	cac His	agt Ser	gcc Ala	tac Tyr	tgc Cys 205	cag Gln	ctt Leu	ttt Phe	624
gtg Val	gtg val 210	gtc Val	ttt Phe	gct Ala	gtc Val	ctt Leu 215	tac Tyr	ttt Phe	ctg Leu	ttg Leu	ccc Pro 220	ctg Leu	ctc Leu	ctc Leu	ata Ile	672
ctt Leu 225	gtg Val	gtc Val	tac Tyr	tgc Cys	agc Ser 230	atg Met	ttc Phe	cga Arg	gtg Val	gcc Ala 235	cgc Arg	gtg Val	gct Ala	gcc Ala	atg Met 240	720
cag Gln	cac His	ggg Gly	ccg Pro	ctg Leu 245	ccc Pro	acg Thr	tgg Trp	atg Met	gag Glu 250	aca Thr	ccc Pro	cgg Arg	caa Gln	cgc Arg 255	tcc Ser	768
gaa Glu	tct Ser	ctc Leu	agc Ser 260	agc Ser	cgc Arg	tcc Ser	acg Thr	atg Met 265	gtc Val	acc Thr	agc Ser	tcg Ser	ggg Gly 270	gcc Ala	ccc Pro	816
cag Gln	acc Thr	acc Thr 275	cca Pro	cac His	cgg Arg	acg Thr	ttt Phe 280	ggg Gly	gga Gly	ggg Gly	aaa Lys	gca Ala 285	gca Ala	gtg Val	gtt val	864
ctc Leu	ctg Leu 290	gct Ala	gtg Val	ggg Gly	gga Gly	cag Gln 295	ttc Phe	ctg Leu	ctc Leu	tgt Cys	tgg Trp 300	ttg Leu	ccc Pro	tac Tyr	ttc Phe	912
tct Ser 305	ttc Phe	cac His	ctc Leu	tat Tyr	gtt Val 310	gcc Ala	ctg Leu	agt Ser	gct Ala	cag Gln 315	ccc Pro	att Ile	tca Ser	act Thr	ggg Gly 320	960
cag Gln	gtg Val	gag Glu	agt Ser	gtg Val 325	gtc Val	acc Thr	tgg Trp	att Ile	ggc Gly 330	tac Tyr	ttt Phe	tgc Cys	ttc Phe	act Thr 335	tcc Ser	1008
aac Asn	cct Pro	ttc Phe	ttc Phe 340	tat Tyr	gga Gly	tgt Cys	ctc Leu	aac Asn 345	cgg Arg	cag Gln	atc Ile	cgg Arg	ggg Gly 350	gag Glu	ctc Leu	1056
	aag Lys															1104
agg Arg	ctg Leu 370	cct Pro	agc Ser	cgg Arg	gag Glu	ggc Gly 375	tcc Ser	att Ile	gag Glu	gag Glu	aac Asn 380	ttc Phe	ctg Leu	cag Gln	ttc Phe	1152
ctt	cag	999	act	ggc	tgt	cct	tct	gag		tgg age		tcc	cga	ccc	cta	1200

```
09/625,191
Leu Gln Gly Thr Gly Cys Pro Ser Glu Ser Trp Val Ser Arg Pro Leu 385 390 395 400
ccc agc ccc aag cag gag cca cct gct gtt gac ttt cga atc cca ggc
                                                                              1248
Pro Ser Pro Lys Gln Glu Pro Pro Ala Val Asp Phe Arg Ile Pro Gly
                  405
                                         410
cag ata gct gag gag acc tct gag ttc ctg gag cag caa ctc acc agc Gln Ile Ala Glu Glu Thr Ser Glu Phe Leu Glu Gln Gln Leu Thr Ser 420 425 430
                                                                              1296
gac atc atc atg tca gac agc tac ctc cgt cct gcc gcc tca ccc cgg
                                                                              1344
Āsp Ile Ile Met Ser Āsp Ser Tyr Leu Arg Pro Āla Āla Ser Pro Arg
ctg gag tca tga
                                                                              1356
Leu Glu Ser
     450
<210>
        451
<211>
<212>
        PRT
<213>
        Homo sapiens
<400>
Met Glu Ser Ser Pro Ile Pro Gln Ser Ser Gly Asn Ser Ser Thr Leu
1 10 15
Gly Arg Val Pro Gln Thr Pro Gly Pro Ser Thr Ala Ser Gly Val Pro
20 25 30
Glu Val Gly Leu Arg Asp Val Ala Ser Glu Ser Val Ala Leu Phe Phe 35 40 45
Met Leu Leu Leu Asp Leu Thr Ala Val Ala Gly Asn Ala Ala Val Met 50 60
Ala Val Ile Ala Lys Thr Pro Ala Leu Arg Lys Phe Val Phe 65 70 75 80
His Leu Cys Leu Val Asp Leu Leu Ala Ala Leu Thr Leu Met Pro Leu
Ala Met Leu Ser Ser Ser Ala Leu Phe Asp His Ala Leu Phe Gly Glu
Val Ala Cys Arg Leu Tyr Leu Phe Leu Ser Val Cys Phe Val Ser Leu
115 120 125
Ala ile Leu Ser Val Ser Ala Ile Asn Val Glu Arg Tyr Tyr Val
    130
```

									09,	/625	.191				
Val 145	His	Pro	Met	Arg	Tyr 150	Glu	val	Arg				Gly	Leu	Val	Ala 160
Ser	val	Leu	Val	Gly 165	Val	Trp	Val	Lys	Ala 170	Leu	Ala	Met	Ala	Ser 175	val
Pro	٧a٦	Leu	Gly 180	Arg	val	Ser	Тгр	Glu 185	Glu	Gly	Ala	Pro	Ser 190	∨al	Pro
Pro	Gly	Cys 195	Ser	Leu	Gln	Тгр	Ser 200	His	Ser	Ala	Tyr	Cys 205	Gln	Leu	Phe
val	val 210	val	Phe	Ala	٧a٦	Leu 215	Tyr	Phe	Leu	Leu	Pro 220	Leu	Leu	Leu	Ile
Leu 225	∨al	٧al	Туг	Cys	Ser 230	Met	Phe	Arg	٧al	Ala 235	Arg	val	Ala	Ala	Met 240
Gln	His	Gly	Pro	Leu 245	Pro	Thr	Trp	Met	G1u 250	Thr	Pro	Arg	Gln	Arg 255	Ser
Glu	Ser	Leu	Ser 260	Ser	Arg	Ser	Thr	меt 265	val	Thr	Ser	Ser	Gly 270	Ala	Pro
Gln	Thr	Thr 275	Pro	ніѕ	Arg	Thr	Phe 280	Gly	Gly	Gly	Lys	Ala 285	Ala	val	∨al
Leu	Leu 290	Ala	val	Gly	Gly	G1n 295	Phe	Leu	Leu	Cys	Trp 300	Leu	Pro	Tyr	Phe
Ser 305	Phe	His	Leu	Tyr	val 310	Ala	Leu	Ser	Ala	Gln 315	Pro	Ile	Ser	Thr	Gly 320
G]n	val	Glu	Ser	va1 325	۷al	Thr	Trp	Ile	Gly 330	туг	Phe	Cys	Phe	Thr 335	Ser
Asn	Pro	Phe	Phe 340	Tyr	Gly	Cys	Leu	Asn 345	Arg	Gln	Ile	Arg	Gly 350	Glu	Leu
Ser	Lys	G]n 355	Phe	Val	Cys	Phe	Phe 360	Lys	Pro	Ala	Pro	G1u 365	Glu	Glu	Leu
Arg	Leu 370	Pro	Ser	Arg	Glu	Gly 375	Ser	Ile	Glu	Glu	Asn 380	Phe	Leu	Gln	Phe
Leu 385	Gln	Gly	Thr	Gly	Cys 390	Pro	Ser	Glu	Ser	Trp 395	val	Ser	Arg	Pro	Leu 400

Pro Ser Pro Lys Gln Glu Pro Pro Ala Val Asp Phe Arg Ile Pro Gly 405 410 415	
Gln Ile Ala Glu Glu Thr Ser Glu Phe Leu Glu Gln Gln Leu Thr Ser 420 425 430	
Asp Ile Ile Met Ser Asp Ser Tyr Leu Arg Pro Ala Ala Ser Pro Arg 435 440 445	
Leu Glu Ser 450	
<210> 3 <211> 969 <212> DNA <213> Homo sapiens	
<220> <221> CDS <222> (1)(969) <223>	
<pre><400> 3 atg gat cca acc atc tca acc ttg gac aca gaa ctg aca cca atc aac Met Asp Pro Thr Ile Ser Thr Leu Asp Thr Glu Leu Thr Pro Ile Asn 1</pre>	48
gga act gag gag act ctt tgc tac aag cag acc ttg agc ctc acg gtg Gly Thr Glu Glu Thr Leu Cys Tyr Lys Gln Thr Leu Ser Leu Thr Val 20 25 30	96
ctg acg tgc atc gtt tcc ctt gtc ggg ctg aca gga aac gca gtt gtg Leu Thr Cys Ile Val Ser Leu Val Gly Leu Thr Gly Asn Ala Val Val 35 40 45	14
ctc tgg ctc ctg ggc tgc cgc atg cgc agg aac gcc ttc tcc atc tac Leu Trp Leu Leu Gly Cys Arg Met Arg Arg Asn Ala Phe Ser Ile Tyr 50 55 60	92
atc ctc aac ttg gcc gca gca gac ttc ctc ttc ctc agc ggc cgc ctt Ile Leu Asn Leu Ala Ala Ala Asp Phe Leu Phe Leu Ser Gly Arg Leu 65 70 75 80	40
ata tat tcc ctg tta agc ttc atc agt atc ccc cat acc atc tct aaa Ile Tyr Ser Leu Leu Ser Phe Ile Ser Ile Pro His Thr Ile Ser Lys 85 90 95	38
atc ctc tat cct gtg atg atg ttt tcc tac ttt gca ggc ctg agc ttt Ile Leu Tyr Pro Val Met Met Phe Ser Tyr Phe Ala Gly Leu Ser Phe 100 105 110	36
ctg agt gcc gtg agc acc gag cgc tgc ctg tcc gtc ctg tgg ccc atc Leu Ser Ala Val Ser Thr Glu Arg Cys Leu Ser Val Leu Trp Pro Ile 115 120 125	34
tgg tac cgc tgc cac cgc ccc aca cac ctg tca gcg gtg gtg tgt gtc Trp Tyr Arg Cys His Arg Pro Thr His Leu Ser Ala Val Val Cys Val 130 135 140 Page 5	32

ctg Leu 145	ctc Leu	tgg Trp	gcc Ala	ctg Leu	tcc Ser 150	ctg Leu	ctg Leu	cgg Arg	agc Ser	atc Ile 155	ctg Leu	gag Glu	tgg Trp	atg Met	tta Leu 160	480
tgt Cys	ggc Gly	ttc Phe	ctg Leu	ttc Phe 165	agt Ser	ggt Gly	gct Ala	gat Asp	tct Ser 170	gct Ala	tgg Trp	tgt Cys	caa Gln	aca Thr 175	tca Ser	528
			aca Thr 180													576
ggg Gly	tcc Ser	agc Ser 195	ctg Leu	gtc val	ctg Leu	ctg Leu	atc Ile 200	agg Arg	att Ile	ctc Leu	tgt Cys	gga Gly 205	tcc Ser	cgg Arg	aag Lys	624
ata Ile	ccg Pro 210	ctg Leu	acc Thr	agg Arg	ctg Leu	tac Tyr 215	gtg val	acc Thr	atc Ile	ctg Leu	ctc Leu 220	aca Thr	gta Val	ctg Leu	gtc Val	672
			tgt Cys													720
tgg Trp	atc Ile	cac His	gtg Val	gac Asp 245	agg Arg	gaa Glu	gtc Val	tta Leu	ttt Phe 250	tgt Cys	cat His	gtt val	cat His	cta Leu 255	gtt Val	768
tct Ser	att Ile	ttc Phe	ctg Leu 260	tcc Ser	gct Ala	ctt Leu	aac Asn	agc Ser 265	agt Ser	gcc Ala	aac Asn	ccc Pro	atc Ile 270	att Ile	tac Tyr	816
ttc Phe	ttc Phe	gtg Val 275	ggc Gly	tcc Ser	ttt Phe	agg Arg	cag Gln 280	cgt Arg	caa Gln	aat Asn	agg Arg	cag Gln 285	aac Asn	ctg Leu	aag Lys	864
			cag Gln													912
gga Gly 305	ggg Gly	cag Gln	ctt Leu	cct Pro	gag Glu 310	gaa Glu	atc Ile	ctg Leu	gag Glu	ctg Leu 315	tcg Ser	gga Gly	agc Ser	aga Arg	ttg Leu 320	960
gag Glu	cag Gln	tga														969
<210 <211 <212	> : !> F	1 322 PRT	sani	ens												

<213> Homo sapiens

<400> 4

Met Asp Pro Thr Ile Ser Thr Leu Asp Thr Glu Leu Thr Pro Ile Asn $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Gly Thr Glu Glu Thr Leu Cys Tyr Lys Gln Thr Leu Ser Leu Thr Val 20 25 30 Page 6 Leu Thr Cys Ile Val Ser Leu Val Gly Leu Thr Gly Asn Ala Val Val 35 40 45

Leu Trp Leu Leu Gly Cys Arg Met Arg Arg Asn Ala Phe Ser Ile Tyr 50 60

Ile Leu Asn Leu Ala Ala Ala Asp Phe Leu Phe Leu Ser Gly Arg Leu 65 70 75 80

Ile Tyr Ser Leu Leu Ser Phe Ile Ser Ile Pro His Thr Ile Ser Lys 85 90 95

Ile Leu Tyr Pro Val Met Met Phe Ser Tyr Phe Ala Gly Leu Ser Phe $100 \hspace{1cm} 105 \hspace{1cm} 110$

Leu Ser Ala Val Ser Thr Glu Arg Cys Leu Ser Val Leu Trp Pro Ile 115 120 125

Trp Tyr Arg Cys His Arg Pro Thr His Leu Ser Ala Val Val Cys Val 130 135 140

Leu Leu Trp Ala Leu Ser Leu Leu Arg Ser Ile Leu Glu Trp Met Leu 145 150 155 160

Cys Gly Phe Leu Phe Ser Gly Ala Asp Ser Ala Trp Cys Gln Thr Ser 165 170 175

Asp Phe Ile Thr Val Ala Trp Leu Ile Phe Leu Cys Val Val Leu Cys 180 185 190

Gly Ser Ser Leu Val Leu Leu Ile Arg Ile Leu Cys Gly Ser Arg Lys 195 200 205

Ile Pro Leu Thr Arg Leu Tyr Val Thr Ile Leu Leu Thr Val Leu Val 210 215 220

Phe Leu Leu Cys Gly Leu Pro Phe Gly Ile Gln Phe Phe Leu Phe Leu 225 230 235 240

Trp Ile His Val Asp Arg Glu Val Leu Phe Cys His Val His Leu Val 245 250 255

Ser Ile Phe Leu Ser Ala Leu Asn Ser Ser Ala Asn Pro Ile Ile Tyr 260 265 270

Phe Phe Val Gly Ser Phe Arg Gln Arg Gln Asn Arg Gln Asn Leu Lys Page 7

275	280	03/023,131	285

Leu Val Leu Gln Arg Ala Leu Gln Asp Ala Ser Glu Val Asp Glu Gly 290 295 300 Gly Gly Gln Leu Pro Glu Glu Ile Leu Glu Leu Ser Gly Ser Arg Leu Glu Gln <210> 5 1017 <211> <212> DNA <213> Homo sapiens <220> <221> CDS <222> <223> (1)..(1017)<400> atg ctg tcc att ttg ctt cct tcc agg gga agc aga agc ggg agc cgt Met Leu Ser Ile Leu Leu Pro Ser Arg Gly Ser Arg Ser Gly Ser Arg 48 cgt gga gct ctg ctc ctg gag gga gcc tcc cgg gac atg gag aag gtg 96 Arg Gly Ala Leu Leu Leu Glu Gly Āla Ser Arg Āsp Met Glu Lys Val 20 gac atg aat aca tca cag gaa caa ggt ctc tgc cag ttc tca gag aag 144 Asp Met Asn Thr Ser Glñ Glu Gln Gly Leu Cŷs Glñ Phe Ser Glū Lys tac aag caa gtc tac ctc tcc ctg gcc tac agt atc atc ttt atc cta 192 Tyr Lys Gln Val Tyr Leu Ser Leu Ala Tyr Ser Ile Ile Phe Ile Leu 55 60 ggg ctg cca cta aat ggc act gtc ttg tgg cac tcc tgg ggc caa acc 240 Gẫy Leu Pro Leu Asn Gẫy Thr Val Leu Trp His Ser Trp Gấy Gln Thr aag cgc tgg agc tgt gcc acc acc tat ctg gtg aac ctg atg gtg gcc Lys Arg Trp Ser Cys Ala Thr Thr Tyr Leu Val Asn Leu Met Val Ala 288 gac ctg ctt tat gtg cta ttg ccc ttc ctc atc atc acc tac tca cta 336 Asp Leu Leu Tyr Val Leu Leu Pro Phe Leu Ile Ile Thr Tyr Ser Leu 105 gat gac agg tgg ccc ttc ggg gag ctg ctc tgc aag ctg gtg cac ttc 384 Asp Asp Arg Trp Pro Phe Gly Glu Leu Leu Cys Lys Leu Val His Phe ctg ttc tat atc aac ctt tac ggc agc atc ctg ctg ctg acc tgc atc Leu Phe Tyr Ile Asn Leu Tyr Gly Ser Ile Leu Leu Leu Thr Cys Ile 432 130

tct gtg cac cag ttc cta ggt gtg tgc cac cca ctg tgt tcg ctg ccc

Page 8

480

Ser V 145	al ı	His	Gln	Phe	Leu 150	Gly	val	Cys	09, His	/625 Pro 155	,191 Leu	Cys	Ser	Leu	Pro 160	
tac c Tyr A	gg : rg	acc Thr	cgc Arg	agg Arg 165	cat His	gcc Ala	tgg Trp	ctg Leu	ggc Gly 170	acc Thr	agc Ser	acc Thr	acc Thr	tgg Trp 175	gcc Ala	528
ctg g Leu V	tg (gtc Val	ctc Leu 180	cag Gln	ctg Leu	ctg Leu	CCC Pro	aca Thr 185	ctg Leu	gcc Ala	ttc Phe	tcc Ser	cac His 190	acg Thr	gac Asp	576
tac a Tyr I	le /	aat Asn 195	ggc Gly	cag Gln	atg Met	atc Ile	tgg Trp 200	tat Tyr	gac Asp	atg Met	acc Thr	agc Ser 205	caa Gln	gag Glu	aat Asn	624
ttt g Phe A 2	at o sp / 10	cgg Arg	ctt Leu	ttt Phe	gcc Ala	tac Tyr 215	ggc Gly	ata Ile	gtt Val	ctg Leu	aca Thr 220	ttg Leu	tct Ser	ggc Gly	ttt Phe	672
ttt c Phe P 225	cc i	tcc Ser	ttg Leu	gtc Val	att Ile 230	ttg Leu	gtg Val	tgc Cys	tat Tyr	tca Ser 235	ctg Leu	atg Met	gtc Val	agg Arg	agc Ser 240	720
ctg a Leu I	tc a le i	aag Lys	cca Pro	gag Glu 245	gag Glu	aac Asn	ctc Leu	atg Met	agg Arg 250	aca Thr	ggc Gly	aac Asn	aca Thr	gcc Ala 255	cga Arg	768
gcc a Ala A	gg 1 rg 9	tcc Ser	atc Ile 260	cgg Arg	acc Thr	atc Ile	cta Leu	ctg Leu 265	gtg Val	tgt Cys	ggc Gly	ctc Leu	ttc Phe 270	acc Thr	ctc Leu	816
tgt t	he 🛚	gtg val 275	ccc Pro	ttc Phe	cat His	atc Ile	act Thr 280	cgc Arg	tcc Ser	ttc Phe	tac Tyr	ctc Leu 285	acc Thr	atc Ile	tgc Cys	864
ttt c Phe Lo 29	tg d eu l 90	ctt Leu	tct Ser	cag Gln	gac Asp	tgc Cys 295	cag Gln	ctc Leu	ttg Leu	atg Met	gca Ala 300	gcc Ala	agt Ser	gtg Val	gcc Ala	912
tac aa Tyr Ly 305	ag a ys I	ata Ile	tgg Trp	agg Arg	cct Pro 310	ctg Leu	gtg Val	agt Ser	gtg Val	agc Ser 315	agc Ser	tgc Cys	ctc Leu	aac Asn	cca Pro 320	960
gtc c Val Lo	tg 1 eu 1	tac Tyr	ttt Phe	ctt Leu 325	tca Ser	agg Arg	ggg Gly	gca Ala	aaa Lys 330	ata Ile	gag Glu	tca Ser	ggc Gly	tcc Ser 335	tcc Ser	1008
aga aa Arg As		tga														1017
<210><211><211><212><213>	PF		sapi	ens												
<400>	6															
Met Le 1	eu S	Ser	Ile	Leu 5	Leu	Pro	Ser	Arg	Gly 10	Ser	Arg	Ser	Gly	Ser 15	Arg	

09/625,191 Arg Gly Ala Leu Leu Glu Gly Ala Ser Arg Asp Met Glu Lys Val 20 25 Asp Met Asn Thr Ser Gln Glu Gln Gly Leu Cys Gln Phe Ser Glu Lys 35 40 Tyr Lys Gln Val Tyr Leu Ser Leu Ala Tyr Ser Ile Ile Phe Ile Leu 50 60 Gly Leu Pro Leu Asn Gly Thr Val Leu Trp His Ser Trp Gly Gln Thr 65 70 75 80 Lys Arg Trp Ser Cys Ala Thr Thr Tyr Leu Val Asn Leu Met Val Ala 85 90 95 Asp Leu Leu Tyr Val Leu Leu Pro Phe Leu Ile Ile Thr Tyr Ser Leu 100 105 110 Asp Asp Arg Trp Pro Phe Gly Glu Leu Leu Cys Lys Leu Val His Phe 115 120 125 Leu Phe Tyr Ile Asn Leu Tyr Gly Ser Ile Leu Leu Leu Thr Cys Ile 130 135 140 Ser Val His Gln Phe Leu Gly Val Cys His Pro Leu Cys Ser Leu Pro 145 150 155 160 Tyr Arg Thr Arg Arg His Ala Trp Leu Gly Thr Ser Thr Thr Trp Ala 165 170 175 Leu Val Val Leu Gln Leu Leu Pro Thr Leu Ala Phe Ser His Thr Asp Tyr Ile Asn Gly Gln Met Ile Trp Tyr Asp Met Thr Ser Gln Glu Asn 195 200 205 Phe Asp Arg Leu Phe Ala Tyr Gly Ile Val Leu Thr Leu Ser Gly Phe 210 220 Phe Pro Ser Leu Val Ile Leu Val Cys Tyr Ser Leu Met Val Arg Ser 225 230 235 240 Leu Ile Lys Pro Glu Glu Asn Leu Met Arg Thr Gly Asn Thr Ala Arg 245 250 255 Ala Arg Ser Ile Arg Thr Ile Leu Leu Val Cys Gly Leu Phe Thr Leu 265 270

											•					
Cy	/s Phe	Val 275	Pro	Phe	His	Ile	Thr 280	Arg	Ser	Phe	Tyr	Leu 285	Thr	Ile	Cys	
Pł	ne Leu 290		Ser	Gln	Asp	Cys 295	Gln	Leu	Leu	Met	Ala 300	Ala	Ser	Val	Ala	
T) 30	r Lys)5	Ile	Тгр	Arg	Pro 310	Leu	val	Ser	٧a٦	Ser 315	Ser	Cys	Leu	Asn	Pro 320	
Vā	ıl Leu	Tyr	Phe	Leu 325	Ser	Arg	Gly	Ala	Lys 330	Ile	Glu	Ser	Gly	Ser 335	Ser	
1Α	g Asn															
<2 <2	211> 212>	7 966 DNA Homo	sap ⁻	iens												
<2 <2		CDS (1).	. (96	6)												
	00> g aac t Asn	7 cag Gln	act Thr	ttg Leu 5	aat Asn	agc Ser	agt Ser	ggg Gly	acc Thr 10	gtg Val	gag Glu	tca Ser	gcc Ala	cta Leu 15	aac Asn	48
t <i>a</i> Ty	t tcc r Ser	aga Arg	ggg G1y 20	agc Ser	aca Thr	gtg Val	cac His	acg Thr 25	gcc Ala	tac Tyr	ctg Le u	gtg Val	ctg Leu 30	agc Ser	tcc Ser	96
ct Le	g gcc u Ala	atg Met 35	ttc Phe	acc Thr	tgc Cys	ctg Leu	tgc Cys 40	ggg Gly	atg Met	gca Ala	ggc Gly	aac Asn 45	agc Ser	atg Met	gtg Val	144
at Il	c tgg e Trp 50	ctg Leu	ctg Leu	ggc Gly	ttt Phe	cga Arg 55	atg Met	cac His	agg Arg	aac Asn	ccc Pro 60	ttc Phe	tgc Cys	atc Ile	tat Tyr	192
at 11 65	c ctc e Leu	aac Asn	ctg Leu	gcg Ala	gca Ala 70	gcc Ala	gac Asp	ctc Leu	ctc Leu	ttc Phe 75	ctc Leu	ttc Phe	agc Ser	atg Met	gct Ala 80	240
t c Se	c acg r Thr	ctc Leu	agc Ser	ctg Leu 85	gaa Glu	acc Thr	cag Gln	ccc Pro	ctg Leu 90	gtc Val	aat Asn	acc Thr	act Thr	gac Asp 95	aag Lys	288
gt Va	c cac l His	gag Glu	ctg Leu 100	atg Met	aag Lys	aga Arg	ctg Leu	atg Met 105	tac Tyr	ttt Phe	gcc Ala	tac Tyr	aca Thr 110	gtg Val	ggc Gly	336
ct Le	g agc u Ser	ctg Leu	ctg Leu	acg Thr	gcc Ala	atc Ile	agc Ser	acc Thr	cag Gln	cgc Arg	tgt Cys	ctc Leu	tct Ser	gtc Val	ctc Leu	384

Page 11

ttc Phe	cct Pro 130	atc Ile	tgg Trp	ttc Phe	aag Lys	tgt Cys 135	cac His	cgg Arg	ccc Pro	agg Arg	cac His 140	ctg Leu	tca Ser	gcc Ala	tgg Trp	432
gtg Val 145	tgt Cys	ggc Gly	ctg Leu	ctg Leu	tgg Trp 150	aca Thr	ctc Leu	tgt Cys	ctc Leu	ctg Leu 155	atg Met	aac Asn	999 Gly	ttg Leu	acc Thr 160	480
tct Ser	tcc Ser	ttc Phe	tgc Cys	agc Ser 165	aag Lys	ttc Phe	ttg Leu	aaa Lys	ttc Phe 170	aat Asn	gaa Glu	gat Asp	cgg Arg	tgc Cys 175	ttc Phe	528
agg Arg	gtg Val	gac Asp	atg Met 180	gtc val	cag Gln	gcc Ala	gcc Ala	ctc Leu 185	atc Ile	atg Met	ggg Gly	gtc val	tta Leu 190	acc Thr	cca Pro	576
gtg Val	atg Met	act Thr 195	ctg Leu	tcc Ser	agc Ser	ctg Leu	acc Thr 200	ctc Leu	ttt Phe	gtc val	tgg Trp	gtg Val 205	cgg Arg	agg Arg	agc Ser	624
	cag Gln 210															672
gcc Ala 225	tct Ser	gtc Val	ctg Leu	gtg Val	ttc Phe 230	ctc Leu	atc Ile	tgt Cys	tcc Ser	ctg Leu 235	cct Pro	ctg Leu	agc Ser	atc Ile	tac Tyr 240	720
tgg Trp	ttt Phe	gtg Val	ctc Leu	tac Tyr 245	tgg Trp	ttg Leu	agc Ser	ctg Leu	ccg Pro 250	ccc Pro	gag Glu	atg Met	cag Gln	gtc Val 255	ctg Leu	768
tgc Cys	ttc Phe	agc Ser	ttg Leu 260	tca Ser	cgc Arg	ctc Leu	tcc Ser	tcg Ser 265	tcc Ser	gta Val	agc Ser	agc Ser	agc Ser 270	gcc Ala	aac Asn	816
	gtc Val															864
acc Thr	agg Arg 290	tcc Ser	ctg Leu	ggg Gly	act Thr	gtg Val 295	ctc Leu	caa Gln	cag Gln	gcg Ala	ctt Leu 300	cgc Arg	gag Glu	gag Glu	ccc Pro	912
gag Glu 305	ctg Leu	gaa Glu	ggt Gly	ggg Gly	gag Glu 310	acg Thr	ccc Pro	acc Thr	gtg Val	ggc Gly 315	acc Thr	aat Asn	gag Glu	atg Met	ggg Gly 320	960
gct Ala	tga															966

<210> 8 <211> 321 <212> PRT <213> Homo sapiens

<400> 8

Met Asn Gln Thr Leu Asn Ser Ser Gly Thr Val Glu Ser Ala Leu Asn $1 \hspace{1cm} 15 \hspace{1cm} 15$ Page 12

Tyr Ser Arg Gly Ser Thr Val His Thr Ala Tyr Leu Val Leu Ser Ser 20 25 30

Leu Ala Met Phe Thr Cys Leu Cys Gly Met Ala Gly Asn Ser Met Val 35 40 45

Ile Trp Leu Leu Gly Phe Arg Met His Arg Asn Pro Phe Cys Ile Tyr 50 60

Ile Leu Asn Leu Ala Ala Ala Asp Leu Leu Phe Leu Phe Ser Met Ala 65 70 75 80

Ser Thr Leu Ser Leu Glu Thr Gln Pro Leu Val Asn Thr Thr Asp Lys 85 90 95

Val His Glu Leu Met Lys Arg Leu Met Tyr Phe Ala Tyr Thr Val Gly
100 105 110

Leu Ser Leu Leu Thr Ala Ile Ser Thr Gln Arg Cys Leu Ser Val Leu 115 120 125

Phe Pro Ile Trp Phe Lys Cys His Arg Pro Arg His Leu Ser Ala Trp 130 135 140

Val Cys Gly Leu Leu Trp Thr Leu Cys Leu Leu Met Asn Gly Leu Thr 145 150 155 160

Ser Ser Phe Cys Ser Lys Phe Leu Lys Phe Asn Glu Asp Arg Cys Phe 165 170 175

Arg Val Asp Met Val Gln Ala Ala Leu Ile Met Gly Val Leu Thr Pro 180 185 190

Val Met Thr Leu Ser Ser Leu Thr Leu Phe Val Trp Val Arg Arg Ser 195 200 205

Ser Gln Gln Trp Arg Arg Gln Pro Thr Arg Leu Phe Val Val Leu 210 215 220

Ala Ser Val Leu Val Phe Leu Ile Cys Ser Leu Pro Leu Ser Ile Tyr 225 230 235 240

Trp Phe Val Leu Tyr Trp Leu Ser Leu Pro Pro Glu Met Gln Val Leu 245 250 255

Cys Phe Ser Leu Ser Arg Leu Ser Ser Ser Val Ser Ser Ser Ala Asn Page 13

265

260

432

Pro Val Ile Tyr Phe Leu Val Gly Ser Arg Arg Ser His Arg Leu Pro 280 Thr Arg Ser Leu Gly Thr Val Leu Gln Gln Ala Leu Arg Glu Glu Pro Glu Leu Glu Gly Gly Glu Thr Pro Thr Val Gly Thr Asn Glu Met Gly Ala <210> 9 <211> 1401 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(1401)<223> <400> atg gaa gct gac ctg ggt gcc act ggc cac agg ccc cgc aca gag ctt Met Glu Ala Asp Leu Gly Ala Thr Gly His Arg Pro Arg Thr Glu Leu 1 5 10 48 gat gat gag gac tcc tac ccc caa ggt ggc tgg gac acg gtc ttc ctg Asp Asp Glu Asp Ser Tyr Pro Gln Gly Gly Trp Asp Thr Val Phe Leu 20 25 3096 gtg gcc ctg ctg ctc ctt ggg ctg cca gcc aat ggg ttg atg gcg tgg 144 Val Ala Leu Leu Leu Gly Leu Pro Ala Asn Gly Leu Met Ala Trp ctg gcc ggc tcc cag gcc cgg cat gga gct ggc acg cgt ctg gcg ctg Leu Ala Gly Ser Gln Ala Arg His Gly Ala Gly Thr Arg Leu Ala Leu 50 55 60 192 ctc ctg ctc agc ctg gcc ctc tct gac ttc ttg ttc ctg gca gcg 240 Leu Leu Leu Ser Leu Ala Leu Ser Asp Phe Leu Phe Leu Ala Ala Ala 70 gcc ttc cag atc cta gag atc cgg cat ggg gga cac tgg ccg ctg ggg Ala Phe Gln Ile Leu Glu Ile Arg His Gly Gly His Trp Pro Leu Gly 288 aca gct gcc tgc cgc ttc tac tac ttc cta tgg ggc gtg tcc tac tcc 336 Thr Ala Ala Cys Arg Phe Tyr Tyr Phe Leu Trp Gly Val Ser Tyr Ser 100 105 tcc ggc ctc ttc ctg ctg gcc gcc ctc agc ctc gac cgc tgc ctg ctg 384 Ser Gly Leu Phe Leu Leu Ala Ala Leu Ser Leu Asp Arg Cys Leu Leu 115 120 125

gcg ctg tgc cca cac tgg tac cct ggg cac cgc cca gtc cgc ctg ccc

Page 14

Ala	Leu 130	Cys	Pro	His	Тгр	Tyr 135	Pro	Gly		/625 Arg		۷al	Arg	Leu	Pro	
ctc Leu 145	tgg Trp	gtc Val	tgc Cys	gcc Ala	ggt Gly 150	gtc val	tgg Trp	gtg Val	ctg Leu	gcc Ala 155	aca Thr	ctc Leu	ttc Phe	agc Ser	gtg Val 160	480
ccc Pro	tgg Trp	ctg Leu	gtc Val	ttc Phe 165	ccc Pro	gag Glu	gct Ala	gcc Ala	gtc Val 170	tgg Trp	tgg Trp	tac Tyr	gac Asp	ctg Leu 175	gtc Val	528
atc Ile	tgc Cys	ctg Leu	gac Asp 180	ttc Phe	tgg Trp	gac Asp	agc Ser	gag Glu 185	gag Glu	ctg Leu	tcg Ser	ctg L eu	agg Arg 190	atg Met	ctg Leu	576
gag Glu	gtc Val	ctg Leu 195	ggg Gly	ggc Gly	ttc Phe	ctg Leu	cct Pro 200	ttc Phe	ctc Leu	ctg Leu	ctg Leu	ctc Leu 205	gtc Val	tgc Cys	cac His	624
gtg Val	ctc Leu 210	acc Thr	cag Gln	gcc Ala	aca Thr	gcc Ala 215	tgt Cys	cgc Arg	acc Thr	tgc Cys	cac His 220	cgc Arg	caa Gln	cag Gln	cag Gln	672
ccc Pro 225	gca Ala	gcc Ala	tgc Cys	cgg Arg	ggc Gly 230	ttc Phe	gcc Ala	cgt Arg	gtg val	gcc Ala 235	agg Arg	acc Thr	att Ile	ctg Leu	tca Ser 240	720
gcc Ala	tat Tyr	gtg Val	gtc val	ctg Leu 245	agg Arg	ctg Leu	ccc Pro	tac Tyr	cag Gln 250	ctg Leu	gcc Ala	cag Gln	ctg Leu	ctc Leu 255	tac Tyr	768
ctg Leu	gcc Ala	ttc Phe	ctg Leu 260	tgg Trp	gac Asp	gtc Val	tac Tyr	tct Ser 265	ggc Gly	tac Tyr	ctg Leu	ctc Leu	tgg Trp 270	gag Glu	gcc Ala	816
ctg Leu	gtc val	tac Tyr 275	tcc Ser	gac Asp	tac Tyr	ctg Leu	atc Ile 280	cta Leu	ctc Leu	aac Asn	agc Ser	tgc Cys 285	ctc Leu	agc Ser	ccc Pro	864
ttc Phe	ctc Leu 290	tgc Cys	ctc Leu	atg Met	gcc Ala	agt Ser 295	gcc Ala	gac Asp	ctc Leu	cgg Arg	acc Thr 300	ctg Leu	ctg Leu	cgc Arg	tcc Ser	912
gtg Val 305	ctc Leu	tcg Ser	tcc Ser	ttc Phe	gcg Ala 310	gca Ala	gct Ala	ctc Leu	tgc Cys	gag Glu 315	gag Glu	cgg Arg	ccg Pro	ggc Gly	agc Ser 320	960
ttc Phe	acg Thr	ccc Pro	act Thr	gag Glu 325	cca Pro	cag Gln	acc Thr	cag Gln	cta Leu 330	gat Asp	tct Ser	gag Glu	ggt Gly	cca Pro 335	act Thr	1008
ctg Leu	cca Pro	gag Glu	ccg Pro 340	atg Met	gca Ala	gag Glu	gcc Ala	cag Gln 345	tca Ser	cag Gln	atg Met	gat Asp	cct Pro 350	gtg Val	gcc Ala	1056
cag Gln	cct Pro	cag Gln 355	gtg Val	aac Asn	ccc Pro	aca Thr	ctc Leu 360	cag Gln	cca Pro	cga Arg	tcg Ser	gat Asp 365	ccc Pro	aca Thr	gct Ala	1104
cag Gln	cca Pro 370	cag Gln	ctg Leu	aac Asn	cct Pro	acg Thr 375	gcc Ala	cag Gln	cca Pro	cag Gln	tcg Ser 380	gat Asp	ccc Pro	aca Thr	gcc Ala	1152

```
09/625,191
cag cca cag ctg aac ctc atg gcc cag cca cag tca gat tct gtg gcc Gln Pro Gln Leu Asn Leu Met Ala Gln Pro Gln Ser Asp Ser Val Ala
                                                                       1200
385
                     390
                                                               400
                                                                       1248
cag cca cag gca gac act aac gtc cag acc cct gca cct gct gcc agt
Gln Pro Gln Ala Asp Thr Asn Val Gln Thr Pro Āla Pro Āla Āla Sēr
                 405
                                      410
                                                           415
tct gtg ccc agt ccc tgt gat gaa gct tcc cca acc cca tcc tcg cat
                                                                       1296
Ser Val Pro Ser Pro Cys Asp Glu Ala Ser Pro Thr Pro Ser Ser His
                                 425
cct acc cca ggg gcc ctt gag gac cca gcc aca cct cct gcc tct gaa
                                                                       1344
Pro Thr Pro Gly Ala Leu Glu Asp Pro Ala Thr Pro Pro Ala Ser Glu
1392
                         455
                                              460
ccc acg tga
                                                                       1401
Pro Thr
465
<210>
       10
<211>
       466
<212>
       PRT
<213>
       Homo sapiens
<400>
       10
Met Glu Ala Asp Leu Gly Ala Thr Gly His Arg Pro Arg Thr Glu Leu
Asp Asp Glu Asp Ser Tyr Pro Gln Gly Gly Trp Asp Thr Val Phe Leu
20 25 30
Val Ala Leu Leu Leu Gly Leu Pro Ala Asn Gly Leu Met Ala Trp
Leu Ala Gly Ser Gln Ala Arg His Gly Ala Gly Thr Arg Leu Ala Leu 50 60
Leu Leu Leu Ser Leu Ala Leu Ser Asp Phe Leu Phe Leu Ala Ala Ala 65 70 75 80
Ala Phe Gln Ile Leu Glu Ile Arg His Gly Gly His Trp Pro Leu Gly
Thr Ala Ala Cys Arg Phe Tyr Tyr Phe Leu Trp Gly Val Ser Tyr Ser
Ser Gly Leu Phe Leu Leu Ala Ala Leu Ser Leu Asp Arg Cys Leu Leu
```

Ala Tyr Val Val Leu Arg Leu Pro Tyr Gln Leu Ala Gln Leu Leu Tyr 245 250 255

Leu Ala Phe Leu Trp Asp Val Tyr Ser Gly Tyr Leu Leu Trp Glu Ala 260 265 270

Leu Val Tyr Ser Asp Tyr Leu Ile Leu Leu Asn Ser Cys Leu Ser Pro 275 280 285

Phe Leu Cys Leu Met Ala Ser Ala Asp Leu Arg Thr Leu Leu Arg Ser 290 295 300

Val Leu Ser Ser Phe Ala Ala Ala Leu Cys Glu Glu Arg Pro Gly Ser 305 310 315 320

Phe Thr Pro Thr Glu Pro Gln Thr Gln Leu Asp Ser Glu Gly Pro Thr 325 330 335

Leu Pro Glu Pro Met Ala Glu Ala Gln Ser Gln Met Asp Pro Val Ala 340 345 350

Gln Pro Gln Val Asn Pro Thr Leu Gln Pro Arg Ser Asp Pro Thr Ala 355 360 365

Gln Pro Gln Leu Asn Pro Thr Ala Gln Pro Gln Ser Asp Pro Thr Ala 370 375 380 Page 17

385	Pro	GIN	Leu	Asn	390	Met	Ala	GIN	Pro	GIN 395	Ser	Asp	Ser	Val	400	
Gln	Pro	Gln	Ala	Asp 405	Thr	Asn	val	Gln	Thr 410	Pro	Ala	Pro	Ala	Ala 415	Ser	
Ser	val	Pro	Ser 420	Pro	Cys	Asp	Glu	Ala 425	Ser	Pro	Thr	Pro	Ser 430	Ser	His	
Pro	Thr	Pro 435	Gly	Ala	Leu	Glu	Asp 440	Pro	Ala	Thr	Pro	Pro 445	Ala	Ser	Glu	
Gly	Glu 450	Ser	Pro	Ser	Ser	Thr 455	Pro	Pro	G∃u	Ala	Ala 460	Pro	Gly	Ala	Gly	
Pro 465	Thr															
<210 <211 <212 <213	L> 9 2> 0	11 993 DNA Homo	sapi	iens												
<220 <221 <222 <223	L> (2> (CDS (1).	. (993	3)												
<40(atg Met 1	gat	l1 cca Pro	acc Thr	acc Thr 5	ccg Pro	gcc Ala	tgg Trp	gga Gly	aca Thr 10	gaa Glu	agt Ser	aca Thr	aca Thr	gtg Val 15	aat Asn	48
gga Gly	aat Asn	gac Asp	caa Gln 20	gcc Ala	ctt Leu	ctt Leu	ctg Leu	ctt Leu 25	tgt Cys	ggc Gly	aag Lys	gag Glu	acc Thr 30	ctg Leu	atc Ile	96
ccg Pro	gtc Val	ttc Phe 35	ctg Leu	atc Ile	ctt Leu	ttc Phe	att Ile 40	gcc Ala	ctg Leu	gtc Val	ggg Gly	ctg Leu 45	gta Val	gga Gly	aac Asn	144
ggg Gly	ttt Phe 50	gtg Val	ctc Leu	tgg Trp	ctc Leu	ctg Leu 55	ggc Gly	ttc Phe	cgc Arg	atg Met	cgc Arg 60	agg Arg	aac Asn	gcc Ala	ttc Phe	192
tct Ser 65	gtc Val	tac Tyr	gtc val	ctc Leu	agc ser 70	ctg Leu	gcc Ala	ggg Gly	gcc Ala	gac Asp 75	ttc Phe	ctc Leu	ttc Phe	ctc Leu	tgc Cys 80	240
ttc Phe	cag Gln	att Ile	ata Ile	aat Asn 85	tgc Cys	ctg Leu	gtg Val	tac Tyr	ctc Leu 90	agt Ser	aac Asn	ttc Phe	ttc Phe	tgt Cys 95	tcc Ser	288
						agc Ser			Thr		va1					336

			100					105	09,	/625	, 191		110			
		gca Ala 115														384
		gtc Val														432
		gcg Ala														480
agc Ser	atc Ile	ttg Leu	gaa Glu	ggg Gly 165	aag Lys	ttc Phe	tgt Cys	ggc Gly	ttc Phe 170	tta Leu	ttt Phe	agt Ser	gat Asp	ggt Gly 175	gac Asp	528
		tgg Trp														576
		ttc Phe 195														624
		tgt Cys														672
atc Ile 225	ctg Leu	ctc Leu	aca Thr	gtg val	ctg Leu 230	gtg val	ttc Phe	ctc Leu	ctc Leu	tgc Cys 235	ggc Gly	ctg Leu	ccc Pro	ttt Phe	ggc Gly 240	720
att Ile	cag Gln	tgg Trp	ttc Phe	cta Leu 245	ata Ile	tta Leu	tgg Trp	atc Ile	tgg Trp 250	aag Lys	gat Asp	tct Ser	gat Asp	gtc Val 255	tta Leu	768
		cat His														816
agt Ser	gcc Ala	aac Asn 275	ccc Pro	atc Ile	att Ile	tac Tyr	ttc Phe 280	ttc Phe	gtg Val	ggc Gly	tct Ser	ttt Phe 285	agg Arg	aag Lys	cag Gln	864
tgg Trp	cgg Arg 290	ctg Leu	cag Gln	cag Gln	ccg Pro	atc Ile 295	ctc Leu	aag Lys	ctg Leu	gct Ala	ctc Leu 300	cag Gln	agg Arg	gct Ala	ctg Leu	912
cag Gln 305	gac Asp	att Ile	gct Ala	gag Glu	gtg Val 310	gat Asp	cac His	agt Ser	gaa Glu	gga Gly 315	tgc Cys	ttc Phe	cgt Arg	cag Gln	ggc Gly 320	960
acc Thr	ccg Pro	gag Glu	atg Met	tcg Ser 325	aga Arg	agc Ser	agt Ser	ctg Leu	gtg Val 330	tag						993
<210 <211 <212 <213	l> : !> !	12 330 PRT Homo	sapi	iens							10					

<400> 12

Met Asp Pro Thr Thr Pro Ala Trp Gly Thr Glu Ser Thr Thr Val Asn $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Gly Asn Asp Gln Ala Leu Leu Leu Leu Cys Gly Lys Glu Thr Leu Ile $20 \hspace{1cm} 25 \hspace{1cm} 30$

Pro Val Phe Leu Ile Leu Phe Ile Ala Leu Val Gly Leu Val Gly Asn 35 40 45

Gly Phe Val Leu Trp Leu Leu Gly Phe Arg Met Arg Arg Asn Ala Phe 50 55 60

Ser Val Tyr Val Leu Ser Leu Ala Gly Ala Asp Phe Leu Phe Leu Cys 70 75 80

Phe Gln Ile Ile Asn Cys Leu Val Tyr Leu Ser Asn Phe Phe Cys Ser 85 90 95

Ile Ser Ile Asn Phe Pro Ser Phe Phe Thr Thr Val Met Thr Cys Ala 100 105 110

Tyr Leu Ala Gly Leu Ser Met Leu Ser Thr Val Ser Thr Glu Arg Cys 115 120 125

Leu Ser Val Leu Trp Pro Ile Trp Tyr Arg Cys Arg Arg Pro Arg His 130 135 140

Leu Ser Ala Val Val Cys Val Leu Leu Trp Ala Leu Ser Leu Leu Leu 145 150 160

Ser Ile Leu Glu Gly Lys Phe Cys Gly Phe Leu Phe Ser Asp Gly Asp 165 170 175

Ser Gly Trp Cys Gln Thr Phe Asp Phe Ile Thr Ala Ala Trp Leu Ile 180 185 190

Phe Leu Phe Met Val Leu Cys Gly Ser Ser Leu Ala Leu Leu Val Arg 195 200 205

Ile Leu Cys Gly Ser Arg Gly Leu Pro Leu Thr Arg Leu Tyr Leu Thr 210 215 220

Ile Leu Leu Thr Val Leu Val Phe Leu Leu Cys Gly Leu Pro Phe Gly 235 240

	Ile	Gln	Trp	Phe	Leu 245	Ile	Leu	Trp	Ile	09, Trp 250	/625 Lys	,191 Asp	Ser	Asp	va1 255	Leu	
I	Phe	Cys	His	Ile 260	ніѕ	Pro	val	Ser	va1 265	val	Leu	Ser	Ser	Leu 270	Asn	Ser	
:	Ser	Ala	Asn 275	Pro	Ile	Ile	Tyr	Phe 280	Phe	∨al	Gly	Ser	Phe 285	Arg	Lys	Gln	
-	Тгр	Arg 290	Leu	Gln	Gln	Pro	Ile 295	Leu	Lys	Leu	Ala	Leu 300	Gln	Arg	Ala	Leu	
•	G]n 305	Asp	Ile	Ala	Glu	va7 310	Asp	His	Ser	Glu	Gly 315	Cys	Phe	Arg	Gln	Gly 320	
-	Thr	Pro	Glu	Met	Ser 325	Arg	Ser	Ser	Leu	va1 330							
	<210 <211 <212 <213	l> : 2> ::	13 1014 DNA Homo	sap ⁻	iens												
	<22(<22(<22(<22(L> (<u>?</u> > (CDS (1).	. (10:	14)												
i	<40(atg Met 1	aat	l3 gag Glu	cca Pro	cta Leu 5	gac Asp	tat Tyr	tta Leu	gca Ala	aat Asn 10	gct Ala	tct Ser	gat Asp	ttc Phe	ccc Pro 15	gat Asp	48
1	tat Tyr	gca Ala	gct Ala	gct Ala 20	ttt Phe	gga Gly	aat Asn	tgc Cys	act Thr 25	gat Asp	gaa Glu	aac Asn	atc Ile	cca Pro 30	ctc Leu	aag Lys	96
í	atg Met	cac His	tac Tyr 35	ctc Leu	cct Pro	gtt Val	att Ile	tat Tyr 40	ggc Gly	att Ile	atc Ile	ttc Phe	ctc Leu 45	gtg Val	gga Gly	ttt Phe	144
F	cca Pro	ggc Gly 50	aat Asn	gca Ala	gta Val	gtg Val	ata Ile 55	tcc Ser	act Thr	tac Tyr	att Ile	ttc Phe 60	aaa Lys	atg Met	aga Arg	cct Pro	192
-	tgg Frp 65	aag Lys	agc Ser	agc Ser	acc Thr	atc Ile 70	att Ile	atg Met	ctg Leu	aac Asn	ctg Leu 75	gcc Ala	tgc Cys	aca Thr	gat Asp	ctg Leu 80	240
l	ctg Leu	tat Tyr	ctg Leu	acc Thr	agc Ser 85	ctc Leu	ccc Pro	ttc Phe	ctg Leu	att Ile 90	cac His	tac Tyr	tat Tyr	gcc Ala	agt Ser 95	ggc Gly	288
							gat Asp										336

ttc Phe	cat His	ttc Phe 115	aac Asn	ctg Leu	tat Tyr	agc Ser	agc Ser 120	atc Ile	ctc	/625 ttc Phe	ctc	acc Thr 125	tgt Cys	ttc Phe	agc Ser	384
atc Ile	ttc Phe 130	cgc Arg	tac Tyr	tgt Cys	gtg Val	atc Ile 135	att Ile	cac His	cca Pro	atg Met	agc Ser 140	tgc Cys	ttt Phe	tcc ser	att Ile	432
cac His 145	aaa Lys	act Thr	cga Arg	tgt Cys	gca Ala 150	gtt val	gta Val	gcc Ala	tgt Cys	gct Ala 155	gtg val	gtg val	tgg Trp	atc Ile	att Ile 160	480
tca Ser	ctg Leu	gta Val	gct Ala	gtc Val 165	att Ile	ccg Pro	atg Met	acc Thr	ttc Phe 170	ttg Leu	atc Ile	aca Thr	tca Ser	acc Thr 175	aac Asn	528
agg Arg	acc Thr	aac Asn	aga Arg 180	tca Ser	gcc Ala	tgt Cys	ctc Leu	gac Asp 185	ctc Leu	acc Thr	agt Ser	tcg Ser	gat Asp 190	gaa Glu	ctc Leu	576
aat Asn	act Thr	att Ile 195	aag Lys	tgg Trp	tac Tyr	aac Asn	ctg Leu 200	att Ile	ttg Leu	act Thr	gca Ala	act Thr 205	act Thr	ttc Phe	tgc Cys	624
ctc Leu	ccc Pro 210	ttg Leu	gtg val	ata Ile	gtg Val	aca Thr 215	ctt Leu	tgc Cys	tat Tyr	acc Thr	acg Thr 220	att Ile	atc Ile	cac His	act Thr	672
ctg Leu 225	acc Thr	cat His	gga Gly	ctg Leu	caa Gln 230	act Thr	gac Asp	agc Ser	tgc Cys	ctt Leu 235	aag Lys	cag Gln	aaa Lys	gca Ala	cga Arg 240	720
agg Arg	cta Leu	acc Thr	att Ile	ctg Leu 245	cta Leu	ctc Leu	ctt Leu	gca Ala	ttt Phe 250	tac Tyr	gta Val	tgt Cys	ttt Phe	tta Leu 255	ccc Pro	768
ttc Phe	cat His	atc Ile	ttg Leu 260	agg Arg	gtc Val	att Ile	cgg Arg	atc Ile 265	gaa Glu	tct Ser	cgc Arg	ctg Leu	ctt Leu 270	tca Ser	atc Ile	816
agt Ser	tgt Cys	tcc Ser 275	att Ile	gag Glu	aat Asn	cag Gln	atc Ile 280	cat His	gaa Glu	gct Ala	tac Tyr	atc Ile 285	gtt Val	tct Ser	aga Arg	864
cca Pro	tta Leu 290	gct Ala	gct Ala	ctg Leu	aac Asn	acc Thr 295	ttt Phe	ggt Gly	aac Asn	ctg Leu	tta Leu 300	cta Leu	tat Tyr	gtg Val	gtg Val	912
gtc Val 305	agc Ser	gac Asp	aac Asn	ttt Phe	cag Gln 310	cag Gln	gct Ala	gtc Val	tgc Cys	tca Ser 315	aca Thr	gtg Val	aga Arg	tgc Cys	aaa Lys 320	960
gta Val	agc Ser	ggg Gly	aac Asn	ctt Leu 325	gag Glu	caa Gln	gca Ala	aag Lys	aaa Lys 330	att Ile	agt Ser	tac Tyr	tca Ser	aac Asn 335	aac Asn	1008
cct Pro	tga															1014

<212> PRT

<213> Homo sapiens

<400> 14

Met Asn Glu Pro Leu Asp Tyr Leu Ala Asn Ala Ser Asp Phe Pro Asp $1 \hspace{1cm} 5 \hspace{1cm} 15$

Tyr Ala Ala Ala Phe Gly Asn Cys Thr Asp Glu Asn Ile Pro Leu Lys 20 25 30

Met His Tyr Leu Pro Val Ile Tyr Gly Ile Ile Phe Leu Val Gly Phe 35 40 45

Pro Gly Asn Ala Val Val Ile Ser Thr Tyr Ile Phe Lys Met Arg Pro 50 60

Trp Lys Ser Ser Thr Ile Ile Met Leu Asn Leu Ala Cys Thr Asp Leu 65 70 75 80

Leu Tyr Leu Thr Ser Leu Pro Phe Leu Ile His Tyr Tyr Ala Ser Gly 85 90 95

Glu Asn Trp Ile Phe Gly Asp Phe Met Cys Lys Phe Ile Arg Phe Ser 100 105 110

Phe His Phe Asn Leu Tyr Ser Ser Ile Leu Phe Leu Thr Cys Phe Ser 115 120 125

Ile Phe Arg Tyr Cys Val Ile Ile His Pro Met Ser Cys Phe Ser Ile 130 135 140

His Lys Thr Arg Cys Ala Val Val Ala Cys Ala Val Val Trp Ile Ile 145 150 155 160

Ser Leu Val Ala Val Ile Pro Met Thr Phe Leu Ile Thr Ser Thr Asn 165 170 175

Arg Thr Asn Arg Ser Ala Cys Leu Asp Leu Thr Ser Ser Asp Glu Leu 180 190

Asn Thr Ile Lys Trp Tyr Asn Leu Ile Leu Thr Ala Thr Thr Phe Cys 195 200 205

Leu Pro Leu Val Ile Val Thr Leu Cys Tyr Thr Thr Ile Ile His Thr 210 220

Leu Thr His Gly Leu Gln Thr Asp Ser Cys Leu Lys Gln Lys Ala Arg 225 230 235 240 Page 23

Arg	Leu	Thr	Ile	Leu 245	Leu	Leu	Leu	Ala	Phe 250	Tyr	val	Cys	Phe	Leu 255	Pro	
Phe	His	Ile	Leu 260	Arg	val	Ile	Arg	Ile 265	Glu	Ser	Arg	Leu	Leu 270	ser	Ile	
Ser	Cys	Ser 275	Ile	Glu	Asn	Gln	11e 280	His	Glu	Ala	Tyr	11e 285	val	Ser	Arg	
Pro	Leu 290	Ala	Ala	Leu	Asn	Thr 295	Phe	Gly	Asn	Leu	Leu 300	Leu	Tyr	val	Val	
va1 305	Ser	Asp	Asn	Phe	Gln 310	Gln	Ala	val	Cys	Ser 315	Thr	val	Arg	Cys	Lys 320	
Val	Ser	Gly	Asn	Leu 325	Glu	Gln	Ala	Lys	Lys 330	Ile	Ser	Tyr	Ser	Asn 335	Asn	
Pro																
<21 <21 <21 <21	1> : 2> :	15 1011 DNA Homo	sap	iens												
<22 <22 <22 <22	1> (2>	CDS (1)	. (10:	11)												
	aac					tgt Cys										48
gct Ala	tta Leu	Pro	atc Ile 20	Ile	Tyr	atc Ile	Leu	Leu	Cys	att Ile	gtt Val	ggt Gly	gtt Val 30	ttt Phe	gga Gly	96
aac Asn	act Thr	ctc Leu 35	tct Ser	caa Gln	tgg Trp	ata Ile	ttt Phe 40	tta Leu	aca Thr	aaa Lys	ata Ile	ggt Gly 45	aaa Lys	aaa Lys	aca Thr	144
tca Ser	acg Thr 50	cac His	atc Ile	tac Tyr	ctg Leu	tca Ser 55	cac His	ctt Leu	gtg Val	act Thr	gca Ala 60	aac Asn	tta Leu	ctt Leu	gtg Val	192
tgc Cys 65	agt Ser	gcc Ala	atg Met	cct Pro	ttc Phe 70	atg Met	agt Ser	atc Ile	tat Tyr	ttc Phe 75	ctg Leu	aaa Lys	ggt Gly	ttc Phe	caa Gln 80	240
tgg Trp	gaa Glu	tat Tyr	caa Gln	tct Ser	gct Ala	caa Gln	tgc Cys	aga Arg	val	gtc Val age	Asn	ttt Phe	ctg Leu	gga Gly	act Thr	288

				85					09, 90	/625	, 191			95		
cta Leu	tcc Ser	atg Met	cat His 100	gca Ala	agt Ser	atg Met	ttt Phe	gtc Val 105	agt Ser	ctc Leu	tta Leu	att Ile	tta Leu 110	agt Ser	tgg Trp	336
att Ile	gcc Ala	ata Ile 115	agc Ser	cgc Arg	tat Tyr	gct Ala	acc Thr 120	tta L eu	atg Met	caa Gln	aag Lys	gat Asp 125	tcc Ser	tcg Ser	caa Gln	384
							aaa Lys									432
aaa Lys 145	ttt Phe	cgc Arg	cag Gln	ccc Pro	aac Asn 150	ttt Phe	gct Ala	aga Arg	aaa Lys	cta Leu 155	tgc Cys	att Ile	tac Tyr	ata Ile	tgg Trp 160	480
gga Gly	gtt Val	gta Val	ctg Leu	ggc Gly 165	ata Ile	atc Ile	att Ile	cca Pro	gtt Val 170	acc Thr	gta Val	tac Tyr	tac Tyr	tca Ser 175	gtc Val	528
ata Ile	gag Glu	gct Ala	aca Thr 180	gaa Glu	gga Gly	gaa Glu	gag Glu	agc Ser 185	cta Leu	tgc Cys	tac Tyr	aat Asn	cgg Arg 190	cag Gln	atg Met	576
gaa Glu	cta Leu	gga Gly 195	gcc Ala	atg Met	atc Ile	tct Ser	cag Gln 200	att Ile	gca Ala	ggt Gly	ctc Leu	att Ile 205	gga Gly	acc Thr	aca Thr	624
ttt Phe	att Ile 210	gga Gly	ttt Phe	tcc Ser	ttt Phe	tta Leu 215	gta Val	gta Val	cta Leu	aca Thr	tca Ser 220	tac Tyr	tac Tyr	tct Ser	ttt Phe	672
gta Val 225	agc Ser	cat His	ctg Leu	aga Arg	aaa Lys 230	ata Ile	aga Arg	acc Thr	tgt Cys	acg Thr 235	tcc Ser	att Ile	atg Met	gag Glu	aaa Lys 240	720
gat Asp	ttg Leu	act Thr	tac Tyr	agt Ser 245	tct Ser	gtg val	aaa Lys	aga Arg	cat His 250	ctt Leu	ttg Leu	gtc Val	atc Ile	cag Gln 255	att Ile	768
cta Leu	cta Leu	ata Ile	gtt Val 260	tgc Cys	ttc Phe	ctt Leu	cct Pro	tat Tyr 265	agt Ser	att Ile	ttt Phe	aaa Lys	ccc Pro 270	att Ile	ttt Phe	816
tat Tyr	gtt Val	cta Leu 275	cac His	caa Gln	aga Arg	gat Asp	aac Asn 280	tgt Cys	cag Gln	caa Gln	ttg Leu	aat Asn 285	tat Tyr	tta Leu	ata Ile	864
gaa Glu	aca Thr 290	aaa Lys	aac Asn	att Ile	ctc Leu	acc Thr 295	tgt Cys	ctt Leu	gct Ala	tcg Ser	gcc Ala 300	aga Arg	agt Ser	agc Ser	aca Thr	912
gac Asp 305	ccc Pro	att Ile	ata Ile	ttt Phe	ctt Leu 310	tta Leu	tta Leu	gat Asp	aaa Lys	aca Thr 315	ttc Phe	aag Lys	aag Lys	aca Thr	cta Leu 320	960
tat Tyr	aat Asn	ctc Leu	ttt Phe	aca Thr 325	aag Lys	tct Ser	aat Asn	tca Ser	gca Ala 330	cat His	atg Met	caa Gln	tca Ser	tat Tyr 335	ggt Gly	1008
tga									P	age	25					1011

Page 25

<210> 16

<211> 336

<212> PRT

<213> Homo sapiens

<400> 16

Met Asn Asn Asn Thr Thr Cys Ile Gln Pro Ser Met Ile Ser Ser Met $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Ala Leu Pro Ile Ile Tyr Ile Leu Leu Cys Ile Val Gly Val Phe Gly 20 25 30

Asn Thr Leu Ser Gln Trp Ile Phe Leu Thr Lys Ile Gly Lys Lys Thr 35 40 45

Ser Thr His Ile Tyr Leu Ser His Leu Val Thr Ala Asn Leu Leu Val 50 55 60

Cys Ser Ala Met Pro Phe Met Ser Ile Tyr Phe Leu Lys Gly Phe Gln 65 70 75 80

Trp Glu Tyr Gln Ser Ala Gln Cys Arg Val Val Asn Phe Leu Gly Thr 85 90 95

Leu Ser Met His Ala Ser Met Phe Val Ser Leu Leu Ile Leu Ser Trp $100 \hspace{1cm} 105 \hspace{1cm} 110$

Ile Ala Ile Ser Arg Tyr Ala Thr Leu Met Gln Lys Asp Ser Ser Gln 115 120 125

Glu Thr Thr Ser Cys Tyr Glu Lys Ile Phe Tyr Gly His Leu Leu Lys 130 135 140

Lys Phe Arg Gln Pro Asn Phe Ala Arg Lys Leu Cys Ile Tyr Ile Trp 145 150 155 160

Gly Val Val Leu Gly Ile Ile Ile Pro Val Thr Val Tyr Tyr Ser Val 165 170 175

Ile Glu Ala Thr Glu Gly Glu Glu Ser Leu Cys Tyr Asn Arg Gln Met 180 185 190

Glu Leu Gly Ala Met Ile Ser Gln Ile Ala Gly Leu Ile Gly Thr Thr 195 200 205

Phe Ile Gly Phe Ser Phe Leu Val Val Leu Thr Ser Tyr Tyr Ser Phe 210 215 220 Page 26

va1 225	Ser	His	Leu	Arg	Lys 230	Ile	Arg	Thr	Cys	Thr 235	Ser	Ile	Met	Glu	Lys 240	
Asp	Leu	Thr	Tyr	Ser 245	Ser	Val	Lys	Arg	нis 250	Leu	Leu	Val	Ile	G]n 255	Ile	
Leu	Leu	Ile	va1 260	Cys	Phe	Leu	Pro	Tyr 265	Ser	Ile	Phe	Lys	Pro 270	Ile	Phe	
Tyr	٧a٦	Leu 275	His	Gln	Arg	Asp	Asn 280	Cys	Gln	Gln	Leu	Asn 285	Tyr	Leu	Ile	
Glu	Thr 290	Lys	Asn	Ile	Leu	Thr 295	Cys	Leu	Ala	Ser	Ala 300	Arg	Ser	Ser	Thr	
Asp 305	Pro	Ile	Ile	Phe	Leu 310	Leu	Leu	Asp	Lys	Thr 315	Phe	Lys	Lys	Thr	Leu 320	
Tyr	Asn	Leu	Phe	Thr 325	Lys	Ser	Asn	Ser	Ala 330	His	Met	Gln	Ser	Tyr 335	Gly	
<210 <211 <212 <213	l>	L7 L041 DNA Homo	sapi	iens												
<220 <221 <222 <223	L> (?> (DS (1).	. (104	11)												
<400 atg Met 1	tac	L7 aac Asn	ggg Gly	tcg Ser 5	tgc Cys	tgc Cys	cgc Arg	atc Ile	gag Glu 10	ggg Gly	gac Asp	acc Thr	atc Ile	tcc Ser 15	cag Gln	48
gtg val	atg Met	Pro	ccg Pro 20	ctg Leu	ctc Leu	att Ile	∨al	gcc Ala 25	ttt Phe	gtg Val	ctg Leu	ggc Gly	gca Ala 30	cta Leu	ggc Gly	96
aat Asn	ggg Gly	gtc Val 35	gcc Ala	ctg Leu	tgt Cys	ggt Gly	ttc Phe 40	tgc Cys	ttc Phe	cac His	atg Met	aag Lys 45	acc Thr	tgg Trp	aag Lys	144
ccc Pro	agc ser 50	act Thr	gtt Val	tac Tyr	ctt Leu	ttc Phe 55	aat Asn	ttg Leu	gcc Ala	gtg Val	gct Ala 60	gat Asp	ttc Phe	ctc Leu	ctt Leu	192
atg Met 65	atc Ile	tgc Cys	ctg Leu	cct Pro	ttt Phe 70	cgg Arg	aca Thr	gac Asp	tat Tyr	tac Tyr 75	ctc Leu	aga Arg	cgt Arg	aga Arg	cac His 80	240
tgg Trp	gct Ala	ttt Phe	ggg Gly	gac Asp	att Ile	ccc Pro	tgc Cys	cga Arg	Val	ggg Gly	Leu	ttc Phe	acg Thr	ttg Leu	gcc Ala	288

85					09, 90	/625	,191			95
ggg	agc	atc	gtg	ttc	ctt	acg	gtg	gtg	gct	gcg

atg Met	aac Asn	agg Arg	gcc Ala 100	ggg Gly	agc Ser	atc Ile	gtg Val	ttc Phe 105	ctt Leu	acg Thr	gtg Val	gtg Val	gct Ala 110	gcg Ala	gac Asp	336
agg Arg	tat Tyr	ttc Phe 115	aaa Lys	gtg Val	gtc Val	cac His	ccc Pro 120	cac His	cac His	gcg Ala	gtg Val	aac Asn 125	act Thr	atc Ile	tcc Ser	384
acc Thr	cgg Arg 130	gtg Val	gcg Ala	gct Ala	ggc Gly	atc Ile 135	gtc Val	tgc Cys	acc Thr	ctg Leu	tgg Trp 140	gcc Ala	ctg Leu	gtc Val	atc Ile	432
ctg Leu 145	gga Gly	aca Thr	gtg Val	tat Tyr	ctt Leu 150	ttg Leu	ctg Leu	gag Glu	aac Asn	cat His 155	ctc Leu	tgc Cys	gtg Val	caa Gln	gag Glu 160	480
acg Thr	gcc Ala	gtc Val	tcc Ser	tgt Cys 165	gag Glu	agc Ser	ttc Phe	atc Ile	atg Met 170	gag Glu	tcg Ser	gcc Ala	aat Asn	ggc Gly 175	tgg Trp	528
cat His	gac Asp	atc Ile	atg Met 180	ttc Phe	cag Gln	ctg Leu	gag Glu	ttc Phe 185	ttt Phe	atg Met	ccc Pro	ctc Leu	ggc Gly 190	atc Ile	atc Ile	576
tta Leu	ttt Phe	tgc Cys 195	tcc Ser	ttc Phe	aag Lys	att Ile	gtt Val 200	tgg Trp	agc Ser	ctg Leu	agg Arg	cgg Arg 205	agg Arg	cag Gln	cag Gln	624
ctg Leu	gcc Ala 210	aga Arg	cag Gln	gct Ala	cgg Arg	atg Met 215	aag Lys	aag Lys	gcg Ala	acc Thr	cgg Arg 220	ttc Phe	atc Ile	atg Met	gtg Val	672
gtg Val 225	gca Ala	att Ile	gtg Val	ttc Phe	atc Ile 230	aca Thr	tgc Cys	tac Tyr	ctg Leu	ccc Pro 235	agc Ser	gtg Val	tct Ser	gct Ala	aga Arg 240	720
ctc Leu	tat Tyr	ttc Phe	ctc Leu	tgg Trp 245	acg Thr	gtg Val	ccc Pro	tcg Ser	agt Ser 250	gcc Ala	tgc Cys	gat Asp	ccc Pro	tct Ser 255	gtc Val	768
cat His	ggg Gly	gcc Ala	ctg Leu 260	cac His	ata Ile	acc Thr	ctc Leu	agc Ser 265	ttc Phe	acc Thr	tac Tyr	atg Met	aac Asn 270	agc Ser	atg Met	816
	gat Asp															864
tac Tyr	aac Asn 290	aag Lys	ctc Leu	aaa Lys	atc Ile	tgc Cys 295	agt Ser	ctg Leu	aaa Lys	ccc Pro	aag Lys 300	cag Gln	cca Pro	gga Gly	cac His	912
tca Ser 305	aaa Lys	aca Thr	caa Gln	agg Arg	ccg Pro 310	gaa Glu	gag Glu	atg Met	cca Pro	att Ile 315	tcg Ser	aac Asn	ctc Leu	ggt Gly	cgc Arg 320	960
agg Arg	agt Ser	tgc Cys	atc Ile	agt Ser 325	gtg Val	gca Ala	aat Asn	agt Ser	ttc Phe 330	caa Gln	agc Ser	cag Gln	tct Ser	gat Asp 335	ggg Gly	1008
caa	tgg	gat	ccc	cac	att	gtt	gag	tgg		tga age :	28					1041

Gln Trp Asp Pro His Ile Val Glu Trp His
340
345

<210> 18

<211> 346

<212> PRT

<213> Homo sapiens

<400> 18

Met Tyr Asn Gly Ser Cys Cys Arg Ile Glu Gly Asp Thr Ile Ser Gln 10 15

Val Met Pro Pro Leu Leu Ile Val Ala Phe Val Leu Gly Ala Leu Gly 20 25 30

Asn Gly Val Ala Leu Cys Gly Phe Cys Phe His Met Lys Thr Trp Lys 35 40 45

Pro Ser Thr Val Tyr Leu Phe Asn Leu Ala Val Ala Asp Phe Leu Leu 50 55 60

Met Ile Cys Leu Pro Phe Arg Thr Asp Tyr Tyr Leu Arg Arg Arg His 65 70 75 80

Trp Ala Phe Gly Asp Ile Pro Cys Arg Val Gly Leu Phe Thr Leu Ala 85 90 95

Met Asn Arg Ala Gly Ser Ile Val Phe Leu Thr Val Val Ala Ala Asp 100 105 110

Arg Tyr Phe Lys Val Val His Pro His His Ala Val Asn Thr Ile Ser 115 120 125

Thr Arg Val Ala Ala Gly Ile Val Cys Thr Leu Trp Ala Leu Val Ile 130 135 140

Leu Gly Thr Val Tyr Leu Leu Leu Glu Asn His Leu Cys Val Gln Glu 145 150 155 160

Thr Ala Val Ser Cys Glu Ser Phe Ile Met Glu Ser Ala Asn Gly Trp 165 170 175

His Asp Ile Met Phe Gln Leu Glu Phe Phe Met Pro Leu Gly Ile Ile 180 185 190

Leu Phe Cys Ser Phe Lys Ile Val Trp Ser Leu Arg Arg Gln Gln 195 200 205

Leu Ala 210		Gln	Ala	Arg	Met 215	Lys	Lys		/625 Thr		Phe	Ile	Met	val	
Val Ala 225	Ile	val	Phe	Ile 230	Thr	Cys	Туг	Leu	Pro 235	Ser	val	Ser	Ala	Arg 240	
Leu Tyr	Phe	Leu	Trp 245	Thr	val	Pro	Ser	ser 250	Ala	Cys	Asp	Pro	Ser 255	val	
His Gly	Ala	Leu 260	ніѕ	Ile	Thr	Leu	Ser 265	Phe	Thr	Tyr	Met	Asn 270	Ser	Met	
Leu Asp	Pro 275	Leu	Val	Tyr	Tyr	Phe 280	Ser	Ser	Pro	Ser	Phe 285	Pro	Lys	Phe	
Tyr Asn 290		Leu	Lys	Ile	Cys 295	Ser	Leu	Lys	Pro	Lys 300	Gln	Pro	Gly	ніѕ	
Ser Lys 305	Thr	Gln	Arg	Pro 310	Glu	Glu	Met	Pro	Ile 315	Ser	Asn	Leu	Gly	Arg 320	
Arg Ser	Cys	Ile	Ser 325	٧a٦	Ala	Asn	Ser	Phe 330	Gln	Ser	Gln	Ser	Asp 335	Gly	
Gln Trp	Asp	Pro 340	ніѕ	Ile	val	Glu	Trp 345	His							
<211> <212>	19 1059 DNA Homo	sap [.]	iens												
	CDS (1).	. (10	59)												
<400> atg ggc Met Gly 1															48
gcc gtg Ala Val	gcg Ala	ctg Leu 20	cta Leu	tcc Ser	aac Asn	gca Ala	ctg Leu 25	gtg Val	ctg Leu	ctt Leu	tgt Cys	tgc Cys 30	gcc Ala	tac Tyr	96
agc gct Ser Ala	gag Glu 35	ctc Leu	cgc Arg	act Thr	cga Arg	gcc Ala 40	tca Ser	ggc Gly	gtc val	ctc Leu	ctg Leu 45	gtg Val	aat Asn	ctg Leu	144
tct ctg Ser Leu 50															192

ctc	ggt Gly	gtg Val	atg Met	cgc	ggg Gly	cgg	aca Thr	ccg	tcg	/625 gcg	ccc	ggc GTv	gca	tgc	caa	24	10
65	diy	VUI	Mec	~ı 9	70	Alg	****	710	361	75	FIU	ч	ΑΙα	Cys	80		
gtc Val	att Ile	ggc Gly	ttc Phe	ctg Leu 85	gac Asp	acc Thr	ttc Phe	ctg Leu	gcg Ala 90	tcc Ser	aac Asn	gcg Ala	gcg Ala	ctg Leu 95	agc Ser	28	38
gtg Val	gcg Ala	gcg Ala	ctg Leu 100	agc Ser	gca Ala	gac Asp	cag Gln	tgg Trp 105	ctg Leu	gca Ala	gtg Val	ggc Gly	ttc Phe 110	cca Pro	ctg Leu	33	6
cgc Arg	tac Tyr	gcc Ala 115	gga Gly	cgc Arg	ctg Leu	cga Arg	ccg Pro 120	cgc Arg	tat Tyr	gcc Ala	ggc Gly	ctg Leu 125	ctg Leu	ctg Leu	ggc Gly	38	34
tgt Cys	gcc Ala 130	tgg Trp	gga Gly	cag Gln	tcg Ser	ctg Leu 135	gcc Ala	ttc Phe	tca Ser	ggc Gly	gct Ala 140	gca Ala	ctt Leu	ggc Gly	tgc Cys	43	2
tcg Ser 145	tgg Trp	ctt Leu	ggc Gly	tac Tyr	agc Ser 150	agc Ser	gcc Ala	ttc Phe	gcg Ala	tcc Ser 155	tgt Cys	tcg Ser	ctg Leu	cgc Arg	ctg Leu 160	48	10
ccg Pro	ccc Pro	gag Glu	cct Pro	gag Glu 165	cgt Arg	ccg Pro	cgc Arg	ttc Phe	gca Ala 170	gcc Ala	ttc Phe	acc Thr	gcc Ala	acg Thr 175	ctc Leu	52	8
cat His	gcc Ala	gtg Val	ggc Gly 180	ttc Phe	gtg Val	ctg Leu	ccg Pro	ctg Leu 185	gcg Ala	gtg Val	ctc Leu	tgc Cys	ctc Leu 190	acc Thr	tcg Ser	57	'6
ctc Leu	cag Gln	gtg Val 195	cac His	cgg Arg	gtg Val	gca Ala	cgc Arg 200	aga Arg	cac His	tgc Cys	cag Gln	cgc Arg 205	atg Met	gac Asp	acc Thr	62	4
gtc Val	acc Thr 210	atg Met	aag Lys	gcg Ala	ctc Leu	gcg Ala 215	ctg Leu	ctc Leu	gcc Ala	gac Asp	ctg Leu 220	cac His	ccc Pro	agg Arg	tat Tyr	67	'2
tgg Trp 225	ccc Pro	agt Ser	gca Ala	tgc Cys	cga Arg 230	cag Gln	gcc Ala	cag Gln	gcc Ala	agg Arg 235	gac Asp	ttg Leu	ggc Gly	gct Ala	ccc Pro 240	72	0
tgg Trp	gca Ala	gtt Val	ggc Gly	ttg Leu 245	agg Arg	agc Ser	ctg Leu	tgg Trp	gca Ala 250	tca Ser	cca Pro	ccg Pro	tta Leu	ctc Leu 255	tgc Cys	76	8
cca Pro	gag Glu	ttc Phe	acc Thr 260	agc Ser	cac His	agc Ser	act Thr	gcc Ala 265	cct Pro	gca Ala	cgc Arg	tgc Cys	tca Ser 270	cag Gln	ggg Gly	81	.6
ttt Phe	cct Pro	gtt Val 275	ggt Gly	tca Ser	ttg Leu	gtg Val	cag Gln 280	aca Thr	ctg Leu	cgg Arg	ggg Gly	cct Pro 285	ctg Leu	cct Pro	cct Pro	86	4
ggg Gly	ata Ile 290	tgt Cys	gct Ala	cac His	agt Ser	gca Ala 295	cag Gln	gga Gly	gct Ala	ttg Leu	cgc Arg 300	aga Arg	gct Ala	gtg Val	ggg Gly	91	2
tgt Cys 305	gct Ala	tct Ser	ccg Pro	gga Gly	ggg Gly 310	gtt Val	ccg Pro	cgg Arg	Āla	ctg Leu 315 age	Leu	tgg Trp	gcg Ala	gcc Ala	aga Arg 320	96	0

									,		,					
cac His	acc Thr	cct Pro	cct Pro	gtg Val 325	cat His	ggc Gly	tgt Cys	ggg Gly	tct Ser 330	gag Glu	gca Ala	tct Ser	gct Ala	tgt Cys 335	ttc Phe	1008
tgc Cys	cca Pro	ctg Leu	ctg Leu 340	acc Thr	cag Gln	tgc Cys	cct Pro	tgc Cys 345	atg Met	gac Asp	ttg Leu	ggc Gly	ttc Phe 350	aag Lys	tct Ser	1056
tga																1059
<210 <211 <211 <211	l> 2>	20 352 PRT Homo	sap ⁻	iens												
<400)>	20														
Met 1	Gly	Pro	Gly	Glu 5	Ala	Leu	Leu	Ala	Gly 10	Leu	Leu	val	Met	val 15	Leu	
Ala	val	Ala	Leu 20	Leu	Ser	Asn	Ala	Leu 25	val	Leu	Leu	Cys	Cys 30	Ala	Tyr	
Ser	Ala	Glu 35	Leu	Arg	Thr	Arg	Ala 40	Ser	Gly	val	Leu	Leu 45	٧al	Asn	Leu	
Ser	Leu 50	Gly	His	Leu	Leu	Leu 55	Ala	Ala	Leu	Asp	Met 60	Pro	Phe	Thr	Leu	
Leu 65	Gly	٧a٦	Met	Arg	G]y 70	Arg	Thr	Pro	Ser	Ala 75	Pro	Gly	Ala	Cys	G]n 80	
۷al	Ile	Gly	Phe	Leu 85	Asp	Thr	Phe	Leu	Ala 90	Ser	Asn	Ala	Ala	Leu 95	Ser	
val	Ala	Ala	Leu 100	Ser	Ala	Asp	Gln	Trp 105	Leu	Ala	val	Gly	Phe 110	Pro	Leu	
Arg	Tyr	Ala 115	Gly	Arg	Leu	Arg	Pro 120	Arg	Tyr	Ala	Gly	Leu 125	Leu	Leu	Gly	
Cys	Ala 130	Trp	Gly	Gln	Ser	Leu 135	Ala	Phe	Ser	Gly	Ala 140	Ala	Leu	Gly	Cys	
Ser 145	Trp	Leu	Gly	Tyr	Ser 150	Ser	Ala	Phe	Ala	Ser 155	Cys	Ser	Leu	Arg	Leu 160	
Pro	Pro	Glu	Pro	Glu 165	Arg	Pro	Arg	Phe	Ala 170	Ala	Phe	Thr	Ala	Thr 175	Leu	

									09	/625	. 191					
His	Ala	Val	Gly 180		val	Leu	Pro	Leu 185	Ala	val	Leu	Cys	Leu 190		Ser	
Leu	Gln	val 195	His	Arg	val	Ala	Arg 200	Arg	His	Cys	Gln	Arg 205	Met	Asp	Thr	
۷al	Thr 210	Met	Lys	Ala	Leu	Ala 215	Leu	Leu	Ala	Asp	Leu 220	His	Pro	Arg	Tyr	
Trp 225	Pro	Ser	Ala	Cys	Arg 230	Gln	Ala	Gln	Ala	Arg 235	Asp	Leu	Gly	Ala	Pro 240	
Trp	Ala	val	Gly	Leu 245	Arg	Ser	Leu	Trp	Ala 250	Ser	Pro	Pro	Leu	Leu 255	Cys	
Pro	Glu	Phe	Thr 260	Ser	ніѕ	Ser	Thr	Ala 265	Pro	Ala	Arg	Cys	Ser 270	Gln	Gly	
Phe	Pro	val 275	Gly	Ser	Leu	val	Gln 280	Thr	Leu	Arg	Gly	Pro 285	Leu	Pro	Pro	
Gly	Ile 290	Cys	Ala	ніѕ	Ser	Ala 295	Gln	Gly	Ala	Leu	Arg 300	Arg	Ala	val	Gly	
Cys 305	Ala	Ser	Pro	Gly	Gly 310	val	Pro	Arg	Ala	Leu 315	Leu	Trp	Ala	Ala	Arg 320	
His	Thr	Pro	Pro	Va1 325	ніѕ	Gly	Cys	Gly	Ser 330	Glu	Ala	Ser	Ala	Cys 335	Phe	
Cys	Pro	Leu	Leu 340	Thr	Gln	Cys	Pro	Cys 345	Met	Asp	Leu	Glу	Phe 350	Lys	Ser	
<210 <211 <212 <213	.> 9 !> [21 984 9NA Homo	sapi	ens												
<220 <221 <222 <223	> (!> (DS (1)	(984	I)												
<400 atg Met 1		21 ccc Pro	aac Asn	agc Ser 5	act Thr	ggc Gly	gag Glu	gtg Val	ccc Pro 10	agc Ser	ccc Pro	att Ile	ccc Pro	aag Lys 15	ggg Gly	48
gct Ala	ttg Leu	ggg Gly	ctc Leu 20	tcc Ser	ctg Leu	gcc Ala	ctg Leu	gca Ala 25	agc Ser	ctc Leu	atc Ile	atc Ile	acc Thr 30	gcg Ala	aac Asn	96

ctg Leu	ctc Leu	cta Leu 35	gcc Ala	ctg Leu	ggc Gly	atc Ile	gct Ala 40	ggg Gly	acc	/625 gcc Ala	acc	tgc Cys 45	gca Ala	gcc Ala	acc Thr	144
tgc Cys	tgg Trp 50	ctg Leu	ctt Leu	ctt Leu	cct Pro	gag Glu 55	cct Pro	act Thr	gct Ala	ggc Gly	tgg Trp 60	gct Ala	gct Ala	cac His	ggg Gly	192
tct Ser 65	ggc Gly	att Ile	gcc Ala	aca Thr	ttg Leu 70	cca Pro	ggg Gly	ctg Leu	tgg Trp	aac Asn 75	cag Gln	agt Ser	cgc Arg	cgg Arg	ggt Gly 80	240
tac Tyr	tgg Trp	tcc Ser	tgc Cys	ctc Leu 85	ctc Leu	gtc Val	tac Tyr	ttg Leu	gct Ala 90	ccc Pro	aac Asn	ttc Phe	tcc Ser	ttc Phe 95	ctc Leu	288
tcc Ser	ctg Leu	ctt Leu	gcc Ala 100	aac Asn	ctc Leu	ttg Leu	ctg Leu	gtg Val 105	cac His	ggg Gly	gag Glu	cgc Arg	tac Tyr 110	atg Met	gca Ala	336
gtc Val	ctg Leu	agg Arg 115	cca Pro	ctc Leu	cag Gln	ccc Pro	cct Pro 120	ggg Gly	agc Ser	att Ile	cgg Arg	ctg Leu 125	gcc Ala	ctg Leu	ctc Leu	384
ctc Leu	acc Thr 130	tgg Trp	gct Ala	ggt Gly	ccc Pro	ctg Leu 135	ctc Leu	ttt Phe	gcc Ala	agt Ser	ctg Leu 140	ccc Pro	gct Ala	ctg Leu	ggg Gly	432
tgg Trp 145	aac Asn	cac His	tgg Trp	acc Thr	cct Pro 150	ggt Gly	gcc Ala	aac Asn	tgc Cys	agc Ser 155	tcc Ser	cag Gln	gct Ala	atc Ile	ttc Phe 160	480
cca Pro	gcc Ala	ccc Pro	tac Tyr	ctg Leu 165	tac Tyr	ctc Leu	gaa Glu	gtc Val	tat Tyr 170	ggg Gly	ctc Leu	ctg Leu	ctg Leu	ccc Pro 175	gcc Ala	528
gtg Val	ggt Gly	gct Ala	gct Ala 180	gcc Ala	ttc Phe	ctc Leu	tct Ser	gtc Val 185	cgc Arg	gtg Val	ctg Leu	gcc Ala	act Thr 190	gcc Ala	cac His	576
cgc Arg	cag Gln	ctg Leu 195	cag Gln	gac Asp	atc Ile	tgc Cys	cgg Arg 200	ctg Leu	gag Glu	cgg Arg	gca Ala	gtg Va1 205	tgc Cys	cgc Arg	gat Asp	624
gag Glu	ccc Pro 210	tcc Ser	gcc Ala	ctg Leu	gcc Ala	cgg Arg 215	gcc Ala	ctt Leu	acc Thr	tgg Trp	agg Arg 220	cag Gln	gca Ala	agg Arg	gca Ala	672
cag Gln 225	gct Ala	gga Gly	gcc Ala	atg Met	ctg Leu 230	ctc Leu	ttc Phe	ggg Gly	ctg Leu	tgc Cys 235	tgg Trp	ggg Gly	ccc Pro	tac Tyr	gtg Val 240	720
gcc Ala	aca Thr	ctg Leu	ctc Leu	ctc Leu 245	tca Ser	gtc Val	ctg Leu	gcc Ala	tat Tyr 250	gag Glu	cag Gln	cgc Arg	ccg Pro	cca Pro 255	ctg Leu	768
ggg Gly	cct Pro	ggg Gly	aca Thr 260	ctg Leu	ttg Leu	tcc Ser	ctc Leu	ctc Leu 265	tcc Ser	cta Leu	gga Gly	agt Ser	gcc Ala 270	agt Ser	gca Ala	816
gcg Ala	gca Ala	gtg Val 275	ccc Pro	gta Val	gcc Ala	atg Met	ggg Gly 280	ctg Leu	Gly	gat Asp age	Gln	cgc Arg 285	tac Tyr	aca Thr	gcc Ala	864

									,		,					
ccc Pro	tgg Trp 290	agg Arg	cag Gln	ccg Pro	ccc Pro	aaa Lys 295	ggt Gly	gcc Ala	tgc Cys	agg Arg	ggc Gly 300	tgt Cys	ggg Gly	gaa Glu	gag Glu	912
cct Pro 305	ccc Pro	ggg Gly	aca Thr	gtc Val	ccg Pro 310	gcc Ala	cca Pro	gca Ala	ttg Leu	cct Pro 315	acc Thr	acc Thr	caa Gln	gca Ala	gcc Ala 320	960
aaa Lys	gca Ala	gtg val	tcg Ser	acc Thr 325	tgg Trp	act Thr	tga									984
<210 <211 <212 <213	L> : 2> 1	22 327 PRT Homo	sap	iens												
<400> 22																
Met 1	Thr	Pro	Asn	Ser 5	Thr	Gly	Glu	Val	Pro 10	Ser	Pro	Ile	Pro	Lys 15	Gly	
Ala	Leu	Gly	Leu 20	Ser	Leu	Ala	Leu	Ala 25	Ser	Leu	Ile	Ile	Thr 30	Ala	Asn	
Leu	Leu	Leu 35	Ala	Leu	Gly	Ile	Ala 40	Glу	Thr	Ala	Ala	Cys 45	Ala	Ala	Thr	
Cys	Trp 50	Leu	Leu	Leu	Pro	Glu 55	Pro	Thr	Ala	Gly	Trp 60	Ala	Ala	нis	Gly	
Ser 65	Gly	Ile	Ala	Thr	Leu 70	Pro	Gly	Leu	Trp	Asn 75	Gln	Ser	Arg	Arg	Gly 80	
Tyr	Тгр	Ser	Cys	Leu 85	Leu	Val	Tyr	Leu	Ala 90	Pro	Asn	Phe	Ser	Phe 95	Leu	
Ser	Leu	Leu	Ala 100	Asn	Leu	Leu	Leu	Val 105	His	Gly	Glu	Arg	Tyr 110	Met	Ala	
٧a٦	Leu	Arg 115	Pro	Leu	Gln	Pro	Pro 120	Gly	Ser	Ile	Arg	Leu 125	Ala	Leu	Leu	
Leu	Thr 130	Тгр	Ala	Gly	Pro	Leu 135	Leu	Phe	Ala	Ser	Leu 140	Pro	Ala	Leu	Gly	
Trp 145	Asn	ніѕ	Тгр	Thr	Pro 150	Gly	Ala	Asn	Cys	Ser 155	Ser	Gln	Ala	Ile	Phe 160	
Pro	Ala	Pro	Tyr	Leu 165	Tyr	Leu	Glu	val	170	Gly age		Leu	Leu	Pro 175	Ala	

val	Gly	Ala	Ala 180	Ala	Phe	Leu	Ser	∨a1 185	Arg	۷al	Leu	Ala	Thr 190	Ala	His	
Arg	Gln	Leu 195	Gln	Asp	Ile	Cys	Arg 200	Leu	Glu	Arg	Ala	va1 205	Cys	Arg	Asp	
Glu	Pro 210	Ser	Ala	Leu	Ala	Arg 215	Ala	Leu	Thr	Trp	Arg 220	Gln	Ala	Arg	Ala	
G]n 225	Ala	Gly	Ala	Met	Leu 230	Leu	Phe	Gly	Leu	Cys 235	Trp	Gly	Pro	Tyr	Val 240	
Ala	Thr	Leu	Leu	Leu 245	Ser	val	Leu	Ala	Tyr 250	Glu	Gln	Arg	Pro	Pro 255	Leu	
Gly	Pro	Gly	Thr 260	Leu	Leu	Ser	Leu	Leu 265	Ser	Leu	Gly	ser	Ala 270	Ser	Ala	
Ala	Ala	va1 275	Pro	val	Ala	Met	Gly 280	Leu	Gly	Asp	Gln	Arg 285	Туг	Thr	Ala	
Pro	Trp 290	Arg	Gln	Pro	Pro	Lys 295	Gly	Ala	Cys	Arg	Gly 300	Cys	Gly	Glu	Glu	
Pro 305	Pro	Gly	Thr	val	Pro 310	Ala	Pro	Ala	Leu	Pro 315	Thr	Thr	Gln	Ala	Ala 320	
Lys	Ala	val	Ser	Thr 325	Тгр	Thr										
<210 <210 <210 <210	1> 1 2> [23 LO44 DNA Homo	sapi	iens												
<220 <221 <221 <221	l> (2> (DS (1)	. (104	14)												
<400 atg Met 1	D> 2 ggg Gly	gat Asp	gag Glu	ctg Leu 5	gca Ala	cct Pro	tgc Cys	cct Pro	gtg Val 10	ggc Gly	act Thr	aca Thr	gct Ala	tgg Trp 15	ccg Pro	48
gcc Ala	ctg Leu	atc Ile	cag G1n 20	ctc Leu	atc Ile	agc Ser	aag Lys	aca Thr 25	ccc Pro	tgc Cys	atg Met	ccc Pro	caa Gln 30	gca Ala	gcc Ala	96
agc Ser	aac Asn	act Thr	tcc Ser	ttg Leu	ggc Gly	ctg Leu	ggg Gly	gac Asp	Leu	agg Arg	val	ccc Pro	agc Ser	tcc Ser	atg Met	144

		33					40					4)				
ctg Leu	tac Tyr 50	tgg Trp	ctt Leu	ttc Phe	ctt Leu	ccc Pro 55	tca Ser	agc Ser	ctg Leu	ctg Leu	gct Ala 60	gca Ala	gcc Ala	aca Thr	ctg Leu	192
gct Ala 65	gtc Val	agc Ser	ccc Pro	ctg Leu	ctg Leu 70	ctg Leu	gtg Val	acc Thr	atc Ile	ctg Leu 75	cgg Arg	aac Asn	caa Gln	cgg Arg	ctg Leu 80	240
cga Arg	cag Gln	gag Glu	ccc Pro	cac His 85	tac Tyr	ctg Leu	ctc Leu	ccg Pro	gct Ala 90	aac Asn	atc Ile	ctg Leu	ctc Leu	tca Ser 95	gac Asp	288
ctg Leu	gcc Ala	tac Tyr	att Ile 100	ctc Leu	ctc Leu	cac His	atg Met	ctc Leu 105	atc Ile	tcc Ser	tcc Ser	agc Ser	agc Ser 110	ctg Leu	ggt Gly	336
ggc Gly	tgg Trp	gag Glu 115	ctg Leu	ggc Gly	cgc Arg	atg Met	gcc Ala 120	tgt Cys	ggc Gly	att Ile	ctc Leu	act Thr 125	gat Asp	gct Ala	gtc Val	384
ttc Phe	gcc Ala 130	gcc Ala	tgc Cys	acc Thr	agc Ser	acc Thr 135	atc Ile	ctg Leu	tcc Ser	ttc Phe	acc Thr 140	gcc Ala	att Ile	gtg Val	ctg Leu	432
cac His 145	acc Thr	tac Tyr	ctg Leu	gca Ala	gtc Val 150	atc Ile	cat His	cca Pro	ctg Leu	cgc Arg 155	tac Tyr	ctc Leu	tcc Ser	ttc Phe	atg Met 160	480
tcc Ser	cat His	ggg Gly	gct Ala	gcc Ala 165	tgg Trp	aag Lys	gca Ala	gtg Val	gcc Ala 170	ctc Leu	atc Ile	tgg Trp	ctg Leu	gtg Val 175	gcc Ala	528
							att Ile									576
cag Gln	ctg Leu	gag Glu 195	gag Glu	caa Gln	gga Gly	gct Ala	tca Ser 200	tac Tyr	atc Ile	cta Leu	cca Pro	cca Pro 205	agc Ser	atg Met	ggc Gly	624
acc Thr	cag Gln 210	ccg Pro	gga Gly	tgt Cys	ggc Gly	ctc Leu 215	ctg Leu	gtc Val	att Ile	gtt Val	acc Thr 220	tac Tyr	acc T h r	tcc Ser	att Ile	672
ctg Leu 225	tgc Cys	gtt Val	ctg Leu	ttc Phe	ctc Leu 230	tgc Cys	aca Thr	gct Ala	ctc Leu	att Ile 235	gcc Ala	aac Asn	tgt Cys	ttc Phe	tgg Trp 240	720
agg Arg	atc Ile	tat Tyr	gca Ala	gag Glu 245	gcc Ala	aag Lys	act T h r	tca Ser	ggc Gly 250	atc Ile	tgg Trp	ggg Gly	cag Gln	ggc Gly 255	tat Tyr	768
tcc Ser	cgg Arg	gcc Ala	agg Arg 260	ggc Gly	acc Thr	ctg Leu	ctg Leu	atc Ile 265	cac His	tca Ser	gtg Val	ctg Leu	atc Ile 270	aca Thr	ttg Le u	816
tac Tyr	gtg Val	agc Ser 275	aca Thr	ggg Gly	gtg Val	gtg Val	ttc Phe 280	tcc Ser	ctg Leu	gac Asp	atg Met	gtg Val 285	ctg Leu	acc Thr	agg Arg	864
tac	cac	cac	att	gac	tct	9 99	act	cac		tgg age		ctg	gca	gct	aac	912

```
09/625,191
Tyr His His Ile Asp Ser Gly Thr His Thr Trp Leu Leu Ala Ala Asn
290 295 300
agt gag gta ctc atg atg ctt ccc cgt gcc atg ctc aca tac ctg tac
Ser Glu Val Leu Met Met Leu Pro Arg Ala Met Leu Thr Tyr Leu Tyr
                                                                                   960
                        310
                                                315
ctg ctc cgc tac cgg cag ctg ttg ggc atg gtc cgg ggc cac ctc cca Leu Leu Arg Tyr Arg Gln Leu Leu Gly Met Val Arg Gly His Leu Pro 325 330 335
                                                                                  1008
tcc agg agg cac cag gcc atc ttt acc att tcc tag
                                                                                  1044
Ser Arg Arg His Gln Ala Ile Phe Thr Ile Ser
<210>
        24
<211>
        347
<212>
        PRT
<213>
        Homo sapiens
<400> 24
Met Gly Asp Glu Leu Ala Pro Cys Pro Val Gly Thr Thr Ala Trp Pro 1 5 10 15
Ala Leu Ile Gln Leu Ile Ser Lys Thr Pro Cys Met Pro Gln Ala Ala
Ser Asn Thr Ser Leu Gly Leu Gly Asp Leu Arg Val Pro Ser Ser Met 35 40
Leu Tyr Trp Leu Phe Leu Pro Ser Ser Leu Leu Ala Ala Ala Thr Leu 50 60
Ala Val Ser Pro Leu Leu Leu Val Thr Ile Leu Arg Asn Gln Arg Leu
Arg Gln Glu Pro His Tyr Leu Leu Pro Ala Asn Ile Leu Leu Ser Asp
Leu Ala Tyr Ile Leu Leu His Met Leu Ile Ser Ser Ser Leu Gly
Gly Trp Glu Leu Gly Arg Met Ala Cys Gly Ile Leu Thr Asp Ala Val
Phe Ala Ala Cys Thr Ser Thr Ile Leu Ser Phe Thr Ala Ile Val Leu
His Thr Tyr Leu Ala Val Ile His Pro Leu Arg Tyr Leu Ser Phe Met
145 150 155 160
```

09/625,191 Ser His Gly Ala Ala Trp Lys Ala Val Ala Leu Ile Trp Leu Val Ala Cys Cys Phe Pro Thr Phe Leu Ile Trp Leu Ser Lys Trp Gln Asp Ala 180 185 190 Gln Leu Glu Gln Gly Ala Ser Tyr Ile Leu Pro Pro Ser Met Gly Thr Gln Pro Gly Cys Gly Leu Leu Val Ile Val Thr Tyr Thr Ser Ile Leu Cys Val Leu Phe Leu Cys Thr Ala Leu Ile Ala Asn Cys Phe Trp Arg Ile Tyr Ala Glu Ala Lys Thr Ser Gly Ile Trp Gly Gln Gly Tyr 245 250 255 Ser Arg Ala Arg Gly Thr Leu Leu Ile His Ser Val Leu Ile Thr Leu Tyr Val Ser Thr Gly Val Val Phe Ser Leu Asp Met Val Leu Thr Arg 280 Tyr His His Ile Asp Ser Gly Thr His Thr Trp Leu Leu Ala Ala Asn 290 295 300 Ser Glu Val Leu Met Met Leu Pro Arg Ala Met Leu Thr Tyr Leu Tyr Leu Leu Arg Tyr Arg Gln Leu Leu Gly Met Val Arg Gly His Leu Pro 325 Ser Arg Arg His Gln Ala Ile Phe Thr Ile Ser <210> 25 1002 <211> <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(1002)<223> <400> 25 ggc ccc cat agg agc caa cga agt cat ctt tgc ttc aga gct aaa cca gly Pro His Arg Ser Gln Arg Ser His Leu Cys Phe Arg Ala Lys Pro

48

gtt Val	ttt Phe	ctt Leu	ctc Leu 20	tcc Ser	aca Thr	gca Ala	aat Asn	atc Ile 25	ttg	/625 aca Thr	gtg	atc Ile	atc Ile 30	ctc Leu	tcc Ser	96
cag Gln																144
ctc Leu	gct Ala 50	gct Ala	gcc Ala	gac Asp	atc Ile	ttg Leu 55	gtc Val	ctc Leu	ttt Phe	ttc Phe	ata Ile 60	gtg val	ttt Phe	gtg Val	gac Asp	192
ttc Phe 65																240
gac Asp	aag Lys	atc Ile	ata Ile	gaa Glu 85	gtg Val	ctg Leu	gaa Glu	ttc Phe	tca Ser 90	tcc Ser	atc Ile	cac His	acc Thr	tcc Ser 95	ata Ile	288
tgg Trp	att Ile	act Thr	gta Val 100	ccg Pro	tta Leu	acc Thr	att Ile	gac Asp 105	agg Arg	tat Tyr	atc Ile	gct Ala	gtc Val 110	tgc Cys	cac His	336
ccg Pro	ctc Leu	aag Lys 115	tac Tyr	cac His	acg Thr	gtc val	tca Ser 120	tac Tyr	cca Pro	gcc Ala	cgc Arg	acc Thr 125	cgg Arg	aaa Lys	gtc Val	384
att Ile	gta Val 130	agt Ser	gtt Val	tac Tyr	atc Ile	acc Thr 135	tgc Cys	ttc Phe	ctg Leu	acc Thr	agc Ser 140	atc Ile	ccc Pro	tat Tyr	tac Tyr	432
tgg Trp 145	tgg Trp	ccc Pro	aac Asn	atc Ile	tgg Trp 150	act Thr	gaa Glu	gac Asp	tac Tyr	atc Ile 155	agc Ser	acc Thr	tct Ser	gtg Val	cat His 160	480
cac His																528
tcc Ser	atc Ile	ttc Phe	ttc Phe 180	atc Ile	ttg Leu	aac Asn	tca Ser	atc Ile 185	att Ile	gtg val	tac Tyr	aag Lys	ctc Leu 190	agg Arg	agg Arg	576
aag Lys	agc Ser	aat Asn 195	ttt Phe	cgt Arg	ctc Leu	cgt Arg	ggc Gly 200	tac Tyr	tcc Ser	acg Thr	ggg Gly	aag Lys 205	acc Thr	acc Thr	gcc Ala	624
atc Ile	ttg Leu 210	ttc Phe	acc Thr	att Ile	acc Thr	tcc Ser 215	atc Ile	ttt Phe	gcc Ala	aca Thr	ctt Leu 220	tgg Trp	gcc Ala	ccc Pro	cgc Arg	672
atc Ile 225	atc Ile	atg Met	att Ile	ctt Leu	tac Tyr 230	cac His	ctc Leu	tat Tyr	ggg Gly	gcg Ala 235	ccc Pro	atc Ile	cag Gln	aac Asn	cgc Arg 240	720
tgg Trp																768
aac Asn									Cys		Ile					816

									95,	023	,					
cgc Arg	acc Thr	atg Met 275	gca Ala	gcc Ala	gcc Ala	acg Thr	ctc Leu 280	aag Lys	gct Ala	ttc Phe	ttc Phe	aag Lys 285	tgc Cys	cag Gln	aag Lys	864
caa Gln	cct Pro 290	gta Val	cag Gln	ttc Phe	tac Tyr	acc Thr 295	aat Asn	cat His	aac Asn	ttt Phe	tcc Ser 300	ata Ile	aca Thr	agt Ser	agc Ser	912
ccc Pro 305	Trp	atc Ile	tcg Ser	ccg Pro	gca Ala 310	aac Asn	tca Ser	cac His	tgc Cys	atc Ile 315	aag Lys	atg Met	ctg Leu	gtg Val	tac Tyr 320	960
cag Gln	tat Tyr	gac Asp	aaa Lys	aat Asn 325	gga Gly	aaa Lys	cct Pro	ata Ile	aaa Lys 330	gta Val	tcc Ser	ccg Pro	tga			1002
<21 <21 <21 <21	1> 2>	26 333 PRT Homo	sap ⁻	i ens												
<40	0>	26														
Gly 1	Pro	His	Arg	ser 5	Gln	Arg	Ser	His	Leu 10	Cys	Phe	Arg	Ala	Lys 15	Pro	
val	Phe	Leu	Leu 20	Ser	Thr	Ala	Asn	Ile 25	Leu	Thr	val	Ile	Ile 30	Leu	Ser	
Gln	Leu	val 35	Ala	Arg	Arg	Gln	Lys 40	Ser	Ser	Tyr	Asn	Tyr 45	Leu	Leu	Ala	
Leu	Аlа 50	Ala	Аlа	Asp	Ile	Leu 55	val	Leu	Phe	Phe	Ile 60	val	Phe	۷al	Asp	
Phe 65	Leu	Leu	Glu	Asp	Phe 70	Ile	Leu	Asn	Met	Gln 75	Met	Pro	Gln	∨al	Pro 80	
Asp	Lys	Ile	Ile	Glu 85	Val	Leu	Glu	Phe	Ser 90	Ser	Ile	ніѕ	Thr	Ser 95	Ile	
Тгр	Ile	Thr	val 100	Pro	Leu	Thr	Ile	Asp 105	Arg	Tyr	Ile	Ala	val 110	Cys	His	
Pro	Leu	Lys 115	Tyr	His	Thr	Val	Ser 120	Tyr	Pro	Ala	Arg	Thr 125	Arg	Lys	Val	
Ile	val 130	Ser	∨al	Tyr	Ile	Thr 135	Cys	Phe	Leu	Thr	Ser 140	Ile	Pro	Tyr	Tyr	
Trp 145	Trp	Pro	Asn	Ile	Trp 150	Thr	Glu	Asp		Ile 155 age		Thr	Ser	val	His 160	

His	Val	Leu	Ile	Trp 165	Ile	His	Cys	Phe	Thr 170	Val	Tyr	Leu	Val	Pro 175	Cys	
Ser	Ile	Phe	Phe 180	Ile	Leu	Asn	Ser	Ile 185	Ile	val	Tyr	Lys	Leu 190	Arg	Arg	
Lys	Ser	Asn 195	Phe	Arg	Leu	Arg	G]y 200	Tyr	Ser	Thr	Gly	Lys 205	Thr	Thr	Ala	
Ile	Leu 210	Phe	Thr	Ile	Thr	Ser 215	Ile	Phe	Ala	Thr	Leu 220	Тгр	Ala	Pro	Arg	
Ile 225	Ile	Met	Ile	Leu	Tyr 230	His	Leu	Туг	Gly	Ala 235	Pro	Ile	Gln	Asn	Arg 240	
Trp	Leu	val	Нis	11e 245	Met	Ser	Asp	Ile	Ala 250	Asn	Met	Leu	Ala	Leu 255	Leu	
Asn	Thr	Ala	Ile 260	Asn	Phe	Phe	Leu	Tyr 265	Cys	Phe	Ile	Ser	Lys 270	Arg	Phe	
Arg	Thr	Met 275	Ala	Ala	Ala	Thr	Leu 280	Lys	Ala	Phe	Phe	Lys 285	Cys	Gln	Lys	
Gln	Pro 290	val	Gln	Phe	туr	Thr 295	Asn	His	Asn	Phe	ser 300	Ile	Thr	Ser	Ser	
Pro 305	Trp	Ile	Ser	Pro	Ala 310	Asn	Ser	His	Cys	Ile 315	Lys	Met	Leu	val	Tyr 320	
Gln	Tyr	Asp	Lys	Asn 325	Gly	Lys	Pro	Ile	Lys 330	val	Ser	Pro				
<210 <211 <212 <213	L> 1 !> [27 L260 DNA Homo	sapi	ens												
<220 <221 <222 <223	> (!> (DS (1)	(126	50)												
<400 atg Met 1	ctg	27 gca Ala	gct Ala	gcc Ala 5	ttt Phe	gca Ala	gac Asp	tct Ser	aac Asn 10	tcc Ser	agc Ser	agc Ser	atg Met	aat Asn 15	gtg Val	48
tcc Ser	ttt Phe	gct Ala	cac His	ctc Leu	cac His	ttt Phe	gcc Ala	gga Gly	Gly	tac Tyr age	Leu	ccc Pro	tct Ser	gat Asp	tcc Ser	96

													•			
cag Gln	gac Asp	tgg Trp 35	aga Arg	acc Thr	atc Ile	atc Ile	ccg Pro 40	gct Ala	ctc Leu	ttg Leu	gtg Val	gct Ala 45	gtc Val	tgc Cys	ctg Leu	144
gtg Val	ggc Gly 50	ttc Phe	gtg Val	gga Gly	aac Asn	ctg Leu 55	tgt Cys	gtg Val	att Ile	ggc Gly	atc Ile 60	ctc Leu	ctt Leu	cac His	aat Asn	192
gct Ala 65	tgg Trp	aaa Lys	gga Gly	aag Lys	cca Pro 70	tcc Ser	atg Met	atc Ile	cac His	tcc Ser 75	ctg Leu	att Ile	ctg Leu	aat Asn	ctc Leu 80	240
agc Ser	ctg Leu	gct Ala	gat Asp	ctc Leu 85	tcc Ser	ctc Leu	ctg Leu	ctg Leu	ttt Phe 90	tct Ser	gca Ala	cct Pro	atc Ile	cga Arg 95	gct Ala	288
acg Thr	gcg Ala	tac Tyr	tcc Ser 100	aaa Lys	agt Ser	gtt Val	tgg Trp	gat Asp 105	cta Leu	ggc Gly	tgg Trp	ttt Phe	gtc Val 110	tgc Cys	aag Lys	336
tcc Ser	tct Ser	gac Asp 115	tgg Trp	ttt Phe	atc Ile	cac His	aca Thr 120	tgc Cys	atg Met	gca Ala	gcc Ala	aag Lys 125	agc Ser	ctg Leu	aca Thr	384
atc Ile	gtt Val 130	gtg val	gtg Val	gcc Ala	aaa Lys	gta Val 135	tgc Cys	ttc Phe	atg Met	tat Tyr	gca Ala 140	agt Ser	gac Asp	cca Pro	gcc Ala	432
aag Lys 145	caa Gln	gtg Val	agt Ser	atc Ile	cac His 150	aac Asn	tac Tyr	acc Thr	atc Ile	tgg Trp 155	tca Ser	gtg Val	ctg Leu	gtg Val	gcc Ala 160	480
atc Ile	tgg Trp	act Thr	gtg Val	gct Ala 165	agc Ser	ctg Leu	tta Leu	ccc Pro	ctg Leu 170	ccg Pro	gaa Glu	tgg Trp	ttc Phe	ttt Phe 175	agc Ser	528
acc Thr	atc Ile	agg Arg	cat His 180	cat His	gaa Glu	ggt Gly	gtg Val	gaa Glu 185	atg Met	tgc Cys	ctc Leu	gtg Val	gat Asp 190	gta Val	cca Pro	576
gct Ala	gtg Val	gct Ala 195	gaa Glu	gag Glu	ttt Phe	atg Met	tcg Ser 200	atg Met	ttt Phe	ggt Gly	aag Lys	ctc Leu 205	tac Tyr	cca Pro	ctc Leu	624
ctg Leu	gca Ala 210	ttt Phe	ggc Gly	ctt Leu	cca Pro	tta Leu 215	ttt Phe	ttt Phe	gcc Ala	agc Ser	ttt Phe 220	tat Tyr	ttc Phe	tgg Trp	aga Arg	672
gct Ala 225	tat Tyr	gac Asp	caa Gln	tgt Cys	aaa Lys 230	aaa Lys	cga Arg	gga Gly	act Thr	aag Lys 235	act Thr	caa Gln	aat Asn	ctt Leu	aga Arg 240	720
aac Asn	cag Gln	ata Ile	cgc Arg	tca Ser 245	aag Lys	caa Gln	gtc Val	aca Thr	gtg Val 250	atg Met	ctg Leu	ctg Leu	agc Ser	att Ile 255	gcc Ala	768
atc Ile	atc Ile	tct Ser	gct Ala 260	ctc Leu	ttg Leu	tgg Trp	ctc Leu	ccc Pro 265	gaa Glu	tgg Trp	gta Val	gct Ala	tgg Trp 270	ctg Leu	tgg Trp	816
gta	tgg	cat	ctg	aag	gct	gca	ggc	ccg		cca age		caa	ggt	ttc	ata	864

Val Trp His Le 275	u Lys Ala	Ala Gly 280	Pro Al	9/625 a Pro	Pro (Gln Gly 285	Phe	Ile	
gcc ctg tct ca Ala Leu Ser Gl 290	a gtc ttg n Val Leu	atg ttt Met Phe 295	tcc at Ser Il	c tct e Ser	tca d	gca aat	cct Pro	ctc Leu	912
att ttt ctt gt Ile Phe Leu Va 305	g atg tcg l Met Ser 310	gaa gag Glu Glu	ttc ag Phe Ar	g gaa g Glu 315	ggc 1 Gly 1	ttg aaa Leu Lys	ggt Gly	gta val 320	960
tgg aaa tgg at Trp Lys Trp Me	g ata acc t Ile Thr 325	aaa aaa Lys Lys	cct cc Pro Pr 33	o Thr	gtc 1 Val 9	tca gag Ser Glu	tct Ser 335	cag Gln	1008
gaa aca cca gc Glu Thr Pro Al 34	a Gly Asn	tca gag Ser Glu	ggt ct Gly Le 345	t cct u Pro	gac a Asp l	aag gtt Lys Val 350	cca Pro	tct Ser	1056
cca gaa tcc cc Pro Glu Ser Pro 355	a gca tcc o Ala Ser	ata cca Ile Pro 360	gaa aa Glu Ly	a gag s Glu	Lys F	ccc agc Pro Ser 365	tct Ser	CCC Pro	1104
tcc tct ggc aa Ser Ser Gly Ly 370	a ggg aaa s Gly Lys	act gag Thr Glu 375	aag gc Lys Al	a gag a Glu	att d Ile F 380	ccc atc Pro Ile	ctt Leu	cct Pro	1152
gac gta gag ca Asp Val Glu Gl 385	g ttt tgg n Phe Trp 390	cat gag ніs Glu	agg ga Arg As	c aca p Thr 395	gtc o	cct tct Pro Ser	gta Val	cag Gln 400	1200
gac aat gac cc Asp Asn Asp Pro	t atc ccc o Ile Pro 405	tgg gaa Trp Glu	cat ga His Gl 41	u Asp	caa g Gln d	gag aca Glu Thr	ggg Gly 415	gaa Glu	1248
ggt gtt aaa ta Gly Val Lys	9								1260
<210> 28 <211> 419 <212> PRT <213> Homo sa	oiens								
<400> 28									
Met Leu Ala Ala 1	a Ala Phe 5	Ala Asp	Ser As 10	n Ser	Ser S	Ser Met	Asn 15	Val	
Ser Phe Ala His 20	s Leu His	Phe Ala	Gly Gly 25	y Tyr	Leu P	Pro Ser 30	Asp	Ser	
Gln Asp Trp Arg	g Thr Ile	Ile Pro 40	Ala Le	ı Leu		Ala Val 15	Cys	Leu	
Val Gly Phe Va	Gly Asn	Leu Cys 55	Val Il	e Gly	Ile L 60	eu Leu	His	Asn	

Ala 65	Trp	Lys	Gly	Lys	Pro 70	Ser	Met	Ile	09, Ніѕ	/625 Ser 75	,191 Leu	Ile	Leu	Asn	Leu 80
Ser	Leu	Ala	Asp	Leu 85	Ser	Leu	Leu	Leu	Phe 90	Ser	Ala	Pro	Ile	Arg 95	Ala
Thr	Ala	Tyr	Ser 100	Lys	Ser	val	Тгр	Asp 105	Leu	Gly	Тгр	Phe	val 110	Cys	Lys
Ser	Ser	Asp 115	Trp	Phe	Ile	His	Thr 120	Cys	Met	Ala	Ala	Lys 125	Ser	Leu	Thr
Ile	Val 130	val	val	Ala	Lys	Val 135	Cys	Phe	Met	Tyr	Ala 140	Ser	Asp	Pro	Ala
Lys 145	Gln	۷al	Ser	Ile	ніs 150	Asn	Tyr	Thr	Ile	Trp 155	Ser	٧a٦	Leu	٧a٦	Ala 160
Ile	Trp	Thr	Val	Ala 165	Ser	Leu	Leu	Pro	Leu 170	Pro	Glu	Trp	Phe	Phe 175	Ser
Thr	Ile	Arg	His 180	His	Glu	Gly	٧a٦	Glu 185	Met	Cys	Leu	۷al	Asp 190	val	Pro
Ala	٧a٦	Ala 195	Glu	Glu	Phe	Met	Ser 200	Met	Phe	Gly	Lys	Leu 205	Tyr	Pro	Leu
Leu	Ala 210	Phe	Gly	Leu	Pro	Leu 215	Phe	Phe	Ala	Ser	Phe 220	Tyr	Phe	Trp	Arg
Ala 225	Tyr	Asp	Gln	Cys	Lys 230	Lys	Arg	Gly	Thr	Lys 235	Thr	Gln	Asn	Leu	Arg 240
Asn	Gln	Ile	Arg	Ser 245	Lys	Gln	val	Thr	va1 250	Met	Leu	Leu	Ser	11e 255	Ala
Ile	Ile	Ser	Ala 260	Leu	Leu	Trp	Leu	Pro 265	Glu	Trp	val	Ala	Trp 270	Leu	Trp
∨al	Trp	ніs 275	Leu	Lys	Ala	Ala	Gly 280	Pro	Ala	Pro	Pro	G]n 285	Gly	Phe	Ile
Ala	Leu 290	Ser	Gln	val	Leu	Met 295	Phe	Ser	Ile	Ser	ser 300	Ala	Asn	Pro	Leu
11e 305	Phe	Leu	val	Met	Ser 310	Glu	Glu	Phe	Arg	Glu 315	Gly	Leu	Lys	Gly	Va1 320

Trp Lys Trp Met Ile Thr Lys Lys Pro Pro Thr Val Ser Glu Ser Gln 325 330 335

Glu Thr Pro Ala Gly Asn Ser Glu Gly Leu Pro Asp Lys Val Pro Ser 340 350

Pro Glu Ser Pro Ala Ser Ile Pro Glu Lys Glu Lys Pro Ser Ser Pro 355 360 365

Ser Ser Gly Lys Gly Lys Thr Glu Lys Ala Glu Ile Pro Ile Leu Pro 370 375 380

Asp Val Glu Gln Phe Trp His Glu Arg Asp Thr Val Pro Ser Val Gln 385 390 395 400

Asp Asn Asp Pro Ile Pro Trp Glu His Glu Asp Gln Glu Thr Gly Glu
405 410 415

Gly Val Lys

<210> 29

<211> 13 <212> PR

<212> PRT <213> Artificial Sequence

<220>

<223> peptide

<400> 29

Arg Arg Leu Ile Glu Asp Ala Glu Tyr Ala Ala Arg Gly 1 10

<210> 30

<211> 11

<212> DNA

<213> Artificial Sequence

<220>

<223> NF kappa B consensus binding site

<400> 30

ggggactttc c

<210> 31

<211> 30 <212> DN/

<212> DNA <213> Artificial Sequence

<220>

<223> primer

11

```
<400> 31
                                                                       30
ccggaattca ccatggatcc aaccaccccg
<210> 32
<211> 31
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 32
ctagtctaga ctctacacca gactgcttct c
                                                                       31
<210> 33
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 33
ttctctgtct acgtcctcag
                                                                       20
<210>
       34
<211>
       20
<212> DNA
<213> Artificial Sequence
<220>
<223> primer
<400> 34
                                                                       20
gtcctgtcat ctcttaacag
<210> 35
<211> 8
<212>
      PRT
<213> Artificial sequence
<220>
<223> Angiopeptin
<220>
<221>
<222>
<223>
       PEPTIDE
       (1)..(7)
<220>
<221>
       MISC_FEATURE
<222>
       (1)..(1)
<223>
       Xaa in position 1 is D-Nal or beta-(1-Naphthyl)-D-alanyl where
       D means the dextrogyre form of the amino acid
<220>
<221> MISC_FEATURE
<222> (4)..(4)
```

```
09/625,191
<223> Xaa in position 4 is D-Trp i.e the dextrogyre form of the amino
        acid, tryptophan
<400> 35
Xaa Cys Tyr Xaa Lys Val Cys Thr
<210> 36
<211>
      8
<212>
      PRT
<213>
      Artificial Sequence
<220>
<223>
       Somatostatin analogue
<220>
      MISC_FEATURE
<221>
<222>
       (4)...(4)
<223>
      Xaa in position 4 is D-Trp i.e the dextrogyre form of the amino
        acid, tryptophan
<220>
<221>
       PEPTIDE
<222>
       (1)..(7)
<223>
<220>
<221>
      MISC_FEATURE
<222>
       (1)..(1)
<223>
      Xaa in position 1 is D2-Nal or beta-(2-Naphthyl)-D-alanyl
       where D means the dextrogyre form of the amino acid
<220>
<221>
      MISC_FEATURE
<222>
       (8)..(8)
<223>
      Xaa in position 1 is D2-Nal or beta-(2-Naphthyl)-D-alanyl
       where D means the dextrogyre form of the amino acid
<400> 36
Xaa Cys Tyr Xaa Lys Val Cys Xaa
1 5
<210>
       37
<211>
      30
<212>
      DNA
<213>
      Artificial Sequence
<220>
<223>
      primer
<400> 37
ccggaattca ccatggagtc ctcacccatc
                                                                       30
```

		09/625,191
<211> <212> <213>	29 DNA Artificial Sequence	03, 023, 232
<220> <223>	primer	
	38 taga catcatgact ccagccggg	29
<212>	39 33 DNA Artificial Sequence	
<220> <223>	primer	
	39 ttca ccatggatcc aaccatctca acc	33
<210> <211> <212> <213>		
<220> <223>	primer	
	40 taga tcactgctcc aatctgcttc	30
<220> <223>	primer	
	41 attc gttatgctgt ccattttgct tcc	33
<210> <211> <212> <213>	42 33 DNA Artificial Sequence	
<220> <223>	primer	
<400> tacttc	42 taga cccaccagca ctcatctgtg tac	33
<210> <211> <212> <213>	43 38 DNA Artificial Sequence	

<220> <223>	primer	
<400> ccggaa	43 ttca ccatgaacca gactttgaat agcagtgg	38
<210> <211> <212> <213>	44 37 DNA Artificial Sequence	
<220> <223>	primer	
<400> ctagtc	44 taga tctcaagccc ccatctcatt ggtgccc	37
<210> <211> <212> <213>		
<220> <223>	primer	
<400> ccggaa	45 ttca ccatggaagc tgacctgg	28
<210> <211> <212> <213>	46 29 DNA Artificial Sequence	
<220> <223>	primer	
<400> ctagtc	46 taga ctcacgtggg gcctgcgcc	29
<210> <211> <212> <213>	47 35 DNA Artificial Sequence	
<220> <223>	primer	
<400> cagagaa	47 attc ctgcaattct attctagctc ctgtg	35
<210> <211> <212> <213>	48 33 DNA Artificial Sequence	
<220> <223>	primer	

<400> gcggga	tcct attgtcaacc aagctgtgac atg	33
<210> <211> <212> <213>	49 27 DNA Artificial Sequence	
<220> <223>	primer	
	49 ttcg ccatgtacaa cgggtcg	27
<210> <211> <212> <213>	50 30 DNA Artificial Sequence	
<220> <223>	primer	
<400> ctagtc	50 taga ttcagtgcca ctcaacaatg	30